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Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 19 05-OCT-2000;
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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                                                                   GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC 300
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synthetic construct
artificial sequences.
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Rance, I., Theisen, M. and Gruber, V.
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/db_xref="taxon:32630"
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AX093047
Gruber, V. and Comeau, D. Synthetic vectors, transgenic for obtaining them
                                      artificial sequences.
1 (bases 1 to 15077)
                                                                synthetic construct
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Gruber, V. and Comeau, D.
                                                                          synthetic construct
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/db_xref="taxon:32630"
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AX036759
AX036759.1
                                                                                                                                                                Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 25 05-OCT-2000;
                                                                                                                                          MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); GRUBER VERONIQUE (FR)
                                                                                                                                                                                                artificial sequences.

1 (bases 1 to 472)

Rance, I., Theisen, M. and Gruber, V.
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/db_xref="taxon:32630"
/note="pMRT1342"
                                                                                            /organism="synthetic construct"
/db_xref="taxon:32630"
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                                                                                                              TGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGA 139
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TTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACCCGGTATGCCGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3
Ax036737
                                                                                                                                                                                                                                                                                                                  Rance,I., Theisen,M. and Gruber,V. Chimeric expression promoters originating i mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 3 05-OCT-2000; MERISTEM THERAPEUTICS (FR); RANCE IANN (FI
                                                                                                                                                                                                                                                                                                                                                                                          synthetic construct. synthetic construct
                                                                                                                                                                                                                                                                                                         MERISTEM THERAPEUTICS (FR);
GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                                      artificial sequences.
1 (bases 1 to 317)
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61 c 74 g 75
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/db_xref="taxon:32630"
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79.98;
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Pred. No. 4.5
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AX036758
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Best Local Similarity
Matches 236; Conserv
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                                                                                                                              GGTTCCCAAGCTTTATTT 335
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                                                                                                                    TGATATCAAGATTGATGT 416
                                                                                                                                                                                                                                                                                                                                                                                                                                               artificial sequences.

1 (bases 1 to 541)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus

Patent: WO 0058485-A 24 05-CCT-2000;
synthetic construct.
synthetic construct
artificial sequences.
                                                 348 bp
Sequence 4 from Patent WO0058485
AX036738
                                        AX036738.1
                                                                    AX036738
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AX036758
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24 from Patent WO0058485.
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/db_xxef="taxon:32630"
/note="promoter MPr1168"
1.541
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91.5%;
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Pred. No. 1.7e-50;
0; Mismatches 21;
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AX036757
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CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAGAAGAA 137
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            al Similarity 99.
212; Conservative
                                                                                                                                                                                                    artificial sequences.

1 (bases 1 to 604)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus

Patent: WO 0058485-A 23 05-OCT-2000;

MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED; GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                                           Sequence
AX036757
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Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 4 05-OCT-2000;
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                                                                                                                                                                                                                                                                                                                      synthetic construct
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                                                        Similarity
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                                                                                                          /note="promoter MPr1167"
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116 c 145 g 15
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/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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70 с
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                                                     49.98;
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99.1%;
                                         0; Mismatches
                                                     Score 196.2; DB 6
Pred. No. 8.1e-46;
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Pred. No. 4.9e-47;
0; Mismatches 1
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138 GATAAGGTCGGTGATTGTGAAAGAGACACATAGAGGACACATGTAAGGTGGAAAATGTAAGG

219 197

CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA

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RESULT 13
AX036741
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                                                                                                                                                       GGTTCCCAAGCTTTATTTCCTTATTTAAGCACTTGTGTAGTAGCTTAGAAAAACCAACACA
                                                                                                                                                                   GGTTCCCAAGCTTTATTTCCTTATTTAAGCACTTGTGTAGTAGCTTAGAAAACCAACACA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
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                                                                                                           ACAACCTAGAGGATCC
                                                                                                                                                                                                                                                    240;
                        Sequence
AX036741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         artificial sequences.

1 (bases 1 to 371)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 5 05-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct.
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/db_xref="taxon:32630"
/note="promoter MPr1146"
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from Patent WO0058485
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synthetic construct
artificial sequences.
1 (bases 1 to 301)
1 (bases 1 to 301)
1 Rance, I., Theisen, M. and Gruber, V.
Rance, I. Theisen, M. and Gruber originating is mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A, 7 05-0CT-2000;
                                                                                                                                                                                                                                                               Rance, I. Theisen, M. and Gruber, V. Chimeric expression promoters originating importate virus and cassava vein mosaic virus Patent: WO 0058485-A 6 05-0CT-2000; MERISTEM THERAPEUTICS (FR); RANCE IANN (FF; GRUBER VERONIQUE (FR)
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/db_xref="taxon:33630"
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/note="promoter MPr1147"
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Pred. No. 8.1e-30;
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Pred. No. 8.1e-30;
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synthetic construct
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1 (bases 1 to 515)
Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 2 05-0CT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR); GRUBER VERONIQUE (FR)
; GRUBER VERONIQUE (FR)
LOCATION/Qualifiers
1 515
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Sequence 2 from Patent WO0058485.
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/db_xref="taxon:32630"
/db_rref="taxon:32630"
/note="Promoter from the intergenic region of Cassava Vein Mosaic Virus of 515 bp in length EMBL U59751"
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1.91
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Pred. No. 5.5e-24;
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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2001B.DAT:*
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ALIGNMENTS

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E Nucleotide sequence of chimeric expression promoter MPr1162.
E Nucleotide sequence of chimeric expression promoter MPr1162.
E Nucleotide sequence of chimeric expression promoter; plant vascular expression promoter; plant vascular expression promoter; plant vascular expression promoter; plant expression promoter; plant expression promoter; Cassava vein mosaic virus;
E Chimeric - Commelina yellow mottle virus.
E Chimeric - Cassava vein mosaic virus.
E CHIMERI - MERISTEM THERAPEUTICS.
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Chimeric
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                                                                                                                                                                                                                                                                                                                                                                                            AAA96854 standard;
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                                           How mottle virus mosaic virus.
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GTGTAGTAGCTTAGAAAACCAACAACAACCTAGAGGATCC
              GTGTAGTAGCTTAGAAAACCAACAACAACAACCTAGAGGATCC
                                                                   ACGCATGCCACGTTACCCGGTATGCCGGTTCCCAAGCTTTATTTCCTTATTTAAGCACTT
                                                                                                                          ACGACTAGTGATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATG
                                                                                                                                                              GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
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                                                                                                           ACGACTAGTGATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATG
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                                                      {	t ACGCATGCCACGTTACCCGGTATGCCGGTTCCCCAAGCTTTATTTTCCTTATTTAAGCACTT}
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Chimeric - Cassava
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GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
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                                          AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGTGGCAT
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60

Query Match Best Local 9 Matches

al Similarity 316; Conserv

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RESULT 4
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                                                      The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
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Chimeric - Cassava
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      intergenic region; Commelina yellow mottle virus; expression promoter; plant vascular expression pro tissue expression promoter; Cassava vein mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Commelina yellow mottle virus Cassava vein mosaic virus.
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                                                The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
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                                                                                     Chimeric expression promoter for transgenic plant production, compris-
sequence from promoter comprising vascular expression region replaced
with sequence from promoter comprising green tissue expression region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTTGGAAAATGTAAGG
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                                                                                                                              2000-647238/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACAACCTAGAGGATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          310;
                                                                                                                                                                                                                                                                                             green tissue expression
genic plant; chimera; ss.
                                                           <u>ن</u>
                                                                                                                                                                                                                                                                                                                 expression promoter;
                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                               Gruber V,
                                                                                                                                                                                                                                                                 Cassava ve
                                                                                                                                                                                                                                                                                                                                             sequence of chimeric expression promoter MPr1116.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                         2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                (first
                                                           81;
                                                                                                                                                                                      99FR-0003925
                                                          91pp;
                                                                                                                                                                                                                                                                 na yellow mottle virus vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                               entry)
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                                                                                                                                                                                                                                                                                                       promoter;
                                                                                                                                                                                                                                                                                                                 plant vascular expression
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                                                                                                                                                                                                                                                                                                        Cassava
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                                                                                                                                                                                                                                                                                                        mosaic
                                                                                                                                                                                                                                                                                                                 promoter;
                                                                                                         comprises
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RESULT 7
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ID AAAS
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                                                                                                                                                                                                                                                                                                                             Chimeric
Chimeric
                                                                                                                                                                                                                                                                                                                                                                                             chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus transgenic plant; chimera; ss.
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  WPI; 2000-647238/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA96858 standard; DNA;
                                              Rance I,
                                                                                                                                          29-MAR-1999;
                                                                                                                                                                                   29-MAR-2000;
                                                                                                                                                                                                                                    05-OCT-2000
                                                                                                                                                                                                                                                                                  WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleotide sequence of chimeric expression promoter MPr1168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGA 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AACCTAGAGGATCC 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTCCCAAGCTTTATTTCCTTATTTAAGCACTTGTGTAGTAGCTTAGAAAACCAACACAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGAAAGTAACCTTATGCATTTGTAACTTG-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC
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                                           Gruber V,
                                                                                                                                                                                                                                                                                                                             Commelina yel
Cassava vein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 107 A; 61 C; 74 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transgenic plants.
                                                                                                                                          99FR-0003925
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79.98;
                                                                                                                                                                                                                                                                                                                             yellow mottle virus ein mosaic virus.
                                           Theisen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                541
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Pred. No. 6e-5
0; Mismatches
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Chimeric expression promoter for transgenic plant production, compris-
sequence from promoter comprising vascular expression region replaced
with sequence from promoter comprising green tissue expression region
                                                                                                                                                                      comprises
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Claim 5; Page 87-88; 91pp; English

The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are producing transgenic plants. useful

Sequence 541 BP; 169 A; 104 C; 130 G; 138 T; 0 other;

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Matches 236;
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399
                     318
                                          339
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                                                                                    279
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                                                                                                                                                  138
                                                                                                                             220
                                                                                                                                                                       160 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 219
                                                                                                                                                                                 78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAA 137
TGATATCAAGATTGATGT
                                        GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197
                    GGTTCCCAAGCTTTATTT 335
                                                                                  GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACGACTAGTGATGATGTGATATCAA
                                                                                                                             GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                        54.0%;
91.5%;
                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                Score 212.4; DB Pred. No. 3e-55; Mismatches
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                                                                                                                                                                                                                 21;
                                                                                                                                                                                                                                      21;
                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                      541;
                                                                                                                                                                                                                1;
                                                                                                                                                                                                               Gaps
                                                              317
                                                                                                                             278
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RESULT 8 AAA96838

AAA96838 standard; DNA; 348 ВP

AAA96838;

19-FEB-2001 (first entry)

Nucleotide sequence of chimeric expression promoter MPr1117

Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression proplant green tissue expression promoter; Cassava vein mosaic transgenic plant; chimera; mosaic promoter;

Chimeric - (Commelina yellow mottle virus Cassava vein mosaic virus.

WO200058485-A1

YER X EXXX 29-MAR-2000; 2000WO-IB00370

29-MAR-1999; 99FR-0003925

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RESULT 9
AAA96857
ID AAA9
AC AAA9
AC AAA9
AC Nucl
XX Prom
KW Prom
KW Chim
KW Plan
KW Plan
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KW W CTAN
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PR 29-P
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Best Local (
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                                                                                                                                                        Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                 Chimeric -
Chimeric -
 29-MAR-1999;
                             29-MAR-2000; 2000WO-IB00370.
                                                           05-OCT-2000
                                                                                     WO200058485-A1
                                                                                                                                                                                                                             Nucleotide sequence of chimeric expression promoter
                                                                                                                                                                                                                                                            19-FEB-2001
                                                                                                                                                                                                                                                                                        AAA96857
                                                                                                                                                                                                                                                                                                                   AAA96857 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 348 BP; 116 A; 70 C; 78 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 5; Page 81; 91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MERI-) MERISTEM THERAPEUTICS
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hes 212;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 264
                                                                                                                                                                                                                                                                                                                                                                                                     GGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
                                                                                                                                                                                                                                                                                                                                                                                                                                             TAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gruber V,
                                                                                                                Commelina yellow mottle virus Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transgenic plants.
  99FR-0003925
                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             51.0%;
99.1%;
                                                                                                                                                                                                                                                                                                                   604
                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 200.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        84 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..2e-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1;
                                                                                                                                                                        vein mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                              MPr1167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             348;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              139
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05-OCT-2000.

WO200058485-A1

29-MAR-2000; 2000WO-IB00370

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RESULT 10
AAA96839
ID AAA966
XX AA96
XX DY 19-FE
XX Promo
KW Promo
KW Chime
KW Plant
KW Plant
KW Plant
KW Plant
KW Chime
OS Chime
OS Chime
PX WO200
XX
XX O5-OC
PD 05-OC
PD 05-OC
PP 29-MA
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Matches
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                                                                                          Chimeric - Chimeric -
                                                                                                                                  Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression proplant green tissue expression promoter; Cassava vein mosaic transgenic plant; chimera; ss.
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The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric expression promoter for transgenic plant production, compris-
sequence from promoter comprising vascular expression region replaced
with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                               19-FEB-2001
                                                                                                                                                                                                                                                          AAA96839
                                                                                                                                                                                                                                                                                   AAA96839 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 604 BP; 186 A; 116 C; 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-647238/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rance
                                                                                                                                                                                                   Nucleotide sequence of chimeric expression promoter MPr1146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       138
                                                                                                                                                                                                                                                                                                                                                     337
                                                                                                                                                                                                                                                                                                                                                                               258
                                                                                                                                                                                                                                                                                                                                                                                                          279
                                                                                                                                                                                                                                                                                                                                                                                                                                   198
                                                                                                                                                                                                                                                                                                                                                                                                                                                            220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hes 221;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present sequence represents a chimeric promoter of the invention.
                                                                                                                                                                                                                                                                                                                                                                   GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGT
                                                                                                                                                                                                                                                                                                                                                                                                                       GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACCT
                                                                                                                                                                                                                                                                                                                                                                                                         5; Page 87; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                          Commelina
Cassava ve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                              (first entry)
                                                                                            vein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                49.98;
                                                                                                        yellow
                                                                                                                                                                                                                                                                                    371
                                                                                            mosaic
                                                                                                       mottle virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                   ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 196.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3e-50;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                               304
                                                                                                                                                              promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                604;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                          336
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RESULT 11
AAA96841
ID AAA96
XX
AC AAA96
XX
ID 19-FE
DT 19-FE
XX
VIOLE
XX
Promc
KW Promc
KW Chime
KW plant
KW trans
XX
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Best Local
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              Promotter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                     Nucleotide sequence of chimeric expression promoter MPr1154
                                                                                                                       19-FEB-2001
                                                                                                                                                                                AAA96841 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 371 BP; 122 A; 68 C; 89 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 5; Page 81; 91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-647238/62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Simes 240;
                                                                                                                                                                                                                                                              356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        producing
                                                                                                                                                                                                                                                                          ACAACCTAGAGGATCC
                                                                                                                                                                                                                                                                                                                  GGTTCCCAAGCTTTATTTCCTTATTTAAGCACTTGTGTAGTAGCTTAGAAAACCAACACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                        GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 189
                                                                                                                                                                                                                                                            ACAACCTAGAGGATCC
                                                                                                                                                                                                                                                                                                                                                                                                             GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACCCGGTATGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                           GCGGAAAGTAACCTTATGCATTTGTAACTTG-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99FR-0003925.
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                                                                                                                                                                                 DNA;
                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          39.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Theisen
                                                                                                                                                                                                                                                                                       393
                                                                                                                                                                                                                                                            371
                                                                                                                                                                                 301
                                                                                                                                                                                 ВP
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Pred. No. 1.3e-37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        371;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    197
                                                                                                                                                                                                                                                                                                                                                  377
                                                                                                                                                                                                                                                                                                                                                                                295
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RESULT 12
AAA96840
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Best Local :
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                                                                  chimeric expression promoter;
plant green tissue expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric expression promoter for transgenic plant production, comprise sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                              Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                      19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric - Commelina yellow mottle virus. Chimeric - Cassava vein mosaic virus.
                                                                                                                       Nucleotide sequence of chimeric expression promoter MPr1147
                                                                                                                                                                                 AAA96840
                                                                                                                                                                                                           AAA96840 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 301 BP; 98 A; 54 C; 74 G; 75 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 5; Page 82; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                182
                                                                                                                                                                                                                                                                                                                                       123
                                                                                                                                                                                                                                                                                                                                                                                              63 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA
                                                                                                                                                                                                                                                                                                                                                                                                                        78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                             GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC
                                                                                                                                                                                                                                                                                                                                      GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000-647238/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99FR-0003925
                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 35.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Theisen
                                                                                                                                                                                                           398
                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 140.8;
Pred. No. 2.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         X
                                                                promoter;
                                                                              plant vascular expression
                                                                  Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .5e-33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      2;
                                                                   vein
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                  mosaic virus;
                                                                              promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                301;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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181 197

Chimeric -Chimeric -

Commelina yellow mottle virus

vein

transgenic plant; chimera;

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                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 153;
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergence regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                Cassava vein mosaic virus;
tissue-specific expression
                                                                                                           Cassava vein
                                                                                                                                CsVMV promoter CVP1.
                                                                                                                                                       18-JUN-1998
                                                                                                                                                                             AAV14019
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAR-1999;
            20-JUN-1997;
                                  24-DEC-1997.
                                                      WO9748819-A1
                                                                                                                                                                                                 AAV14019 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a chimeric promoter of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200058485-A1
                                                                                                                                                                                                                                                                                                                       138
                                                                                                                                                                                                                                                                                                                                             160
                                                                                                                                                                                                                                                        279
                                                                                                                                                                                                                                                                                                220 GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                     78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                      producing transgenic plants.
                                                                                                                                                                                                                                                                  GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
                                                                                                                                                                                                                                                                                                               GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                           CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000-647238/62
                                                                                                                                                                                                                                                       GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
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                                                                           vein
                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 82; 91pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                  398 BP; 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99FR-0003925
            97WO-US10376
                                                                                                                                                                                                 DNA; 392
                                                                            virus
                                                                                                                                                                                                                                                                                                                                                                                               35.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                A; 80 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Theisen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                               CSVMV; promoter; cultivated crop; control; transgenic plant; ss.
                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                     Score 140.8; DB 2
Pred. No. 2.8e-33;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                93 G;
                                                                                                                                                                                                                                                                                                                                                                                                                               97 T; 0 other;
                                                                                                                                                                                                                                                       314
                                                                                                                                                                                                                                                                                                                                                                                                          DB 21; Length
                                                                                                                                                                                                                                                                                                                                                                                      2
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                                                                                                                                                                                                                                                                                                                                                                                                            398;
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RESULT 14
AAV14021
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Best Local S
Matches 126
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                                                                                                                                                                                                                                              Cassava vein mosaic virus; CsVMV; promoter; cultivated tissue-specific expression control; transgenic plant; s
                                                                                                                                                                                                                                                                                           CsvMV promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 74; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cassava vein mosaic virus promoter -
DNA sequences for producing transgeni
WPI; 1998-063157/06
                            Beachy RN,
                                                                                           20-JUN-1996;
                                                                                                                        20-JUN-1997;
                                                                                                                                                      24 - DEC - 1997
                                                                                                                                                                                   WO9748819-A1
                                                                                                                                                                                                                   Cassava
                                                                                                                                                                                                                                                                                                                           18-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                       AAV14021 standard; DNA; 411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 phenotype.
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                                                           (SCRI ) SCRIPPS RES INST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93 GAATATTGAAGACGTAAGCACTGACCACAACAATGAAAAGAAGAAGAAGATAAGGTCGGTGAT 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAATCTTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT
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                            De Kochko A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        De Kochko A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 154 A;
                                                                                                                                                                                                                   mosaic
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93.3%;
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                              Fauquet C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fauquet
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transgenic plants
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Pred. No. 4.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  83 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     red. No. 4.5e
Mismatches
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                              Verdaguer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.5e-27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
          Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a cassava vein mosaic virus promoter, and is a nucleic acid molecule of the invention. The promoter is capable of initiating transcription of an operably linked heterologous nucleic acid sequence in a plant cell. The CSVMV promoters are active in both monocot and dicot plant species, and therefore can be readily applied to a variety of cultivated crops. Although generally constitutive, the derivative promoters include promoters that can regulate expression in a tissue-specific manner, and therefore are useful for controlling expression of heterologous genes in a tissue-specific manner. The promoters can be used for producing transgenic plants with an altered
                                                                                                                                                                                                                                                           Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                    plant
                                                                                                                                                                                                                                                                                                             Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter;
                                                                                                                                                                                                                                                                                                                                                                                     19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA96836 standard; DNA; 515 BP
                                                                                                                                                                                                                               WO200058485-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cassava vein mosaic virus
DNA sequences for producir
                                                                                                                    (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                         29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                      transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         phenotype(s)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATGCATTTGTAATTT 227
                                                                                        I,
                                                                                                                                                                                                                                                                                                green tissue expression promoter; Cassava
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 76-77; 115pp; English.
                                                                                        Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 157 A; 60 C;
                                                                                                                                                                                                                                                                                                                                                        intergenic region of Cassava vein mosaic virus
                                                                                                                                               99FR-0003925.
                                                                                                                                                                                                                                                                                      SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ic virus promoter - used to express heterologous
producing transgenic plants having altered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30.7%;
        comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                87 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  107 T;
        green
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -27;
9;
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       expression region
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                                                                           Best Local Sin
Matches 126;
                                                                                                                 Query Match
                                                                                                                                                                                                                           The present sequence represents a promoter fragment from the intergenic region of Cassava vein mosalc virus. The promoter is used to construct chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the
                                                                                                                                                     Sequence 515 BP; 198 A; 79 C; 109 G; 129 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 80;
 216
                   93 GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAAGATAAGGTCGGTGAT 152
                                                                                                                                                                                         ond plant promoter originates from the Cassava vein mosaic virus. chimeric promoters are useful for producing transgenic plants.
 GAATCTTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAAGATAAGGTCGGTGAT
                                                                                               Similarity
                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     91pp; English
                                                                                           30.7%;
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                                                                                             Score 120.6;
Pred. No. 5e-
                                                                           Mismatches
                                                                                               5e-27;
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Search completed: April 14, 2003, 08:19:36 Job time: 129.948 secs

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ATGCATTTGTAATTT 227

212 335

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ATCACAAAGGAATCT

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393
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       Issued_Patents_NA: *
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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/cgn2_6/ptodata/1/ina/backfiles1.seq:*
                           US-08-232-463-14
US-08-247-809A-3
US-08-711-728-3
US-08-764-100-23
US-09-91-238-1
US-09-330-760-1
US-09-330-737-1
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US-09-330-734A-5
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Query Match
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ALIGNMENTS	US-07-966-187-17	US-08-711-728-5	US-08-247-809A-5	US-09-065-999-6	US-09-065-999-5	US-09-458-093-2	US-08-894-440-2	US-09-471-913-6	US-09-185-244-2	US-08-729-601A-46	US-09-011-151-9	US-09-011-151-8	US-09-363-970-35	US-08-729-601A-43	US-07-936-163-46	5254799-30	US-08-463-308-30	US-08-446-486-30
	Sequence 17, Appl	Sequence 5, Appli	Sequence 5, Appli	Sequence 6, Appli	Sequence 5, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 6, Appli	״	Sequence 46, Appl	Sequence 9, Appli	Sequence 8, Appli	Sequence 35, Appl	Sequence 43, Appl	Sequence 46, Appl	Patent No. 5254799	•	Sequence 30, Appl

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; IMMEDIATE SOURCE:
CLONE: pTZgpt-F1s
US-08-232-463-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       : Sequence 14, Application US/08232463
: Patent No. 5670367
: GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-232-463-14/c
                                                                                                 TELEFAX: (703)663-4109
TELEX: 899149
INFORMATION FOR SEQ ID NO: 14
SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: DORNER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKMER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
                                                                                                                                                                                                                                                                      APPLICATION NUMBER: EP 91 114 300.6 FILING DATE: 26-AUG-1991 ATTORNEY/AGENT INFORMATION: NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768 REFERENCE/DOCKET NUMBER: 30472/114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (703)836-9300
TELEFAX: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 1800 Diagonal Road, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                 STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION: 435
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STATE: VA
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Length 7218;

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                              Query Match
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Matches 18; Conserv
   Matches
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MEDIUM TYPE: Diskette, 3.50 inch

MEDIUM TYPE: storage

COMPUTER: NEC POWERMATE 1 Plus

OPERATING SYSTEM: DOS

SPTWARE: WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Peter
APPLICANT: Edgar
                                                                                                                 INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 439 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
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CORRESPONDENCE ADDRESS:
ADDRESSEE: SDBTWG ....
                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: Bayer 9049-KGB TELECOMMUNICATION INFORMATION: TELEPHONE: (914) 332-1700
                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: P 43 178 45.6 (Germany)
FILING DATE: May 28, 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1058 CTCCCTCGACCTGCAGCCAAGCTCGGAA 1031
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                Local
                                                                                                     STRANDEDNESS:
                                                                                                                                                                                            TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                       NAME: Kurt G. Briscoe
REGISTRATION NUMBER:
 39;
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                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New York
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                                                                                                                                                                                            (914) 332-1844
9.9%;
ilarity 100.0%;
Conservative
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Edgar Maiss
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                                                                         DNA (genomic)
                                                                                                      single
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               Score 39;
Pred. No.
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RESULT 3
US-08-711-728-3
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                                                                                                                                                                   Sequence 23, Application US/08764100 Patent No. 5773700
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APPLICANT: Peter H. Scl
APPLICANT: Edgar Maiss
                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 33,141
REFERENCE/DOCKET NUMBER: BATELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER
FILING PARTY NUMBERS N
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STREET: New York
STATE: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 28-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Kurt G. Briscoe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
MEDIUM TYPE: storage
                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                       APPLICANT:
                                                                      APPLICANT:
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       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                    338 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 376
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OPERATING SYSTEM: DOS
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39; Conserv
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van Grinsven J., Martinus
De Haan, Petrus T.
Gielen L., Johannes J.
Peters, Dirk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (914) 332-1700
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; Pred. No.
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Goldbach, Robert W

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US-09-042-426-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/032,235
FILING DATE: 17-MAR-1993
APPLICATION NUMBER: GB 9206016.9
FILING DATE: 19-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO. 577370071s, Allen E.
REGISTRATION NUMBER: 34,490
REFERENCE/DOCKET NUMBERD: 137-1061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 857-1125 INFORMATION FOR SEQ ID NO:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/042,426
                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6114608artis Corporation
STREET: 564 Morris Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 446 base pairs
TYPE: nucleic acid
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                        TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION: (415)_354-3592
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APPLICATION NUMBER: (
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                                                                                                                                                                                                                                                                                                                                                                                                                                      274 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 312
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ZIP: 943
                                                                                                                                                             STREET: 564 Morris
CITY: Summit
STATE: New Jersey
                                                                                                                                   COUNTRY: UZIP: 07901
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                             INFORMATION:
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                                                                                                                                                                                                                                                        Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
VENTION: DNA Construct Containing Bacillus
EQUENCES: 11
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Pred. No.
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Best Local
                                                                           TELEFAX: (919) 541-868
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                 REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
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NAME: HOXIE, Thomas
REGISTRATION UNMBER: 32,993
REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                            APPLICATION NUMBER: 09/042,426
FILING DATE: March 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,93
                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/291,238
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ADDRESSEE: No. 6222104artis Corporation
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LENGTH: 532 base pairs
TYPE: nucleic acid
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IMMEDIATE SOURCE:
CLONE: 35S Promoter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
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MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                             TYPE: nucleic acid
STRANDEDNESS: doub
                 TOPOLOGY:
                                                                                                                               TELEPHONE:
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                                                              LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Summit: New Jersey
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                                                                532 base pairs
                                                                                                                 (919) 541-8689
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DNA (genomic)
                                 double
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DNA Construct Containing Bacillus
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100.0%; Pred. No.
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               RESULT 8
US-09-328-473-1
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Sequence
                                                                                                                                 Matches
                                                                                                                                             Query Match
Best Local (
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                           NAME: HOX1e, Thomas
REGISTRAE: 32,993
REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: DNA CONSTRUCT CONTAIN NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6229075artis Corporation
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CLONE: 35S Promoter
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CLONE: 35S Promoter
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APPLICATION NUMBER: 09/0.
FILING DATE: March 13, 1:
ATTORNEY/AGENT INFORMATION:
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les 39; Conservative
                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: double
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CLASSIFICATION:
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1, Application US/09328473
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ilarity 100.0%;
Conservative
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100.0%; Pr
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DNA Construct Containing Bacillus
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13, 1998
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Pred. No.
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Pred. No.
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0.0079;
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US-09-330-737-1
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Best Local S
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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MOLECULE TYPE: DNA
HYPOTHETICAL: NO
HYPOTHETICAL: NO
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               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION: (919) 541-8614
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PRIOR APPLICATION NUMBER:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                              NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                   APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
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CLONE: 35S Promoter
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ATTORNEY/AGENT INFORMATION:
NAME: Hoxie, Thomas
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CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6232533art1s Corporation
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                                                                                                                   STATE:
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SOFTWARE:
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New Jersey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    532 base pairs
                                                                                                                                                 564 Morris Avenue
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 PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                 No. 6232534artis Corporation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 4;
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/330,737

FILING DATE

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US-09-329-169-1
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REFERENCE/DOCKET NUMBER: 135/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/09329169 Patent No. 6329575 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 9.9%; Score 39; DB Best Local Similarity 100.0%; Pred. No. 0.1 Matches 39; Conservative 0; Mismatches
               TELEFAX: (919) 5
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09
FILING DATE: March 13, 19
ATTORNEY/AGENT INFORMATION:
NAME: Hoxie, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMMEDIATE SOURCE:
CLONE: 35S Promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL:
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: doub
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                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/329,169
FILING DATE: 09-Jun-1999
CLASSIFICATION: <UNHOWN>
PRIOR APPLICATION DATA:
                                                                    REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
                                                                                                                               APPLICATION NUMBER: 09/042,426
ETLING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Hoxie, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 07901
                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Summit
                                                  TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: No. 6329575artis Corporation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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13, 1998
                                   541-8689
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                                                                                  ; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-330-714A-1
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GENERAL INFORMATION:
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                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 1
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                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 09/042,426
FILING DATE: <Unknown>
ATTORNBY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION INFORMATION:
/ Match 9.9%; Score 39; DB 4; Length 532; Local Similarity 100.0%; Pred. No. 0.0079; ndels 39; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible operating SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                           ANTI-SENSE: NO
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 07901
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: 35S Promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
Thuringiensis Gene
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                                                                                                                                                                                    HYPOTHETICAL: NO
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                                                                                                                                                                                                                                           LENGTH: 532 base pairs TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/330,714A FILING DATE: 11-Jun-1999 CLASSIFICATION: <Unknown>
                                                                                                                          CLONE:
                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (91)
TELEFAX: (919)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Summit
STATE: New Jersey
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                                                                                                                          35S Promoter
                                                                                                                                                                                                                                                                                                                                                                  (919)
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                                                                                                                                                                                                                                                                                                                                                 541-8689
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Pred. No.
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258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296

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RESULT 13
US-09-042-426-5
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US-09-328-826-1
                                                                                                 Sequence 5, Application US/09042426
Patent No. 6114608
GENERAL INFORMATION:
APPLICANT: Irvin J. Mettler, Pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09328826
Patent No. 6399860
Patent No. 6399860
GENERAL INFORMATION:
APPLICANT: IIVIN J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
              APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCE: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6114608artis Corporation
                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPHONE: (919) 541-8689
    STREET:
                                                                                                                                                                                                                                                                                                                                                                          Local
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS: ADDRESSEE: No. 6399860art1s Corporation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 312
                                                                                                                                                                                                                                                                                                                                                ch 9.9%; Sc
l Similarity 100.0%; F
39; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 532 base pairs TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/328,826 FILING DATE: 09-Jun-1999 CLASSIFICATION: <Unknown>
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/042,426 FILING DATE: March 13, 1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 07901
564 Morris Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           35S Promoter
                                                                                                                                                                                                                                                                                                                                                                       Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                    Mismatches
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 560 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best
                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
ATTORNEY/AGENT INFORMATION:
NAME: HOXLE, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/042,426
FILING DATE: March 13, 1998
                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Hoxie, Thomas
REGISTRATION NUMBER: 32,993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                             STREET: JUNE STREET: Summit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/0 FILING DATE: March 13, 1998 CLASSIFICATION: 800
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                                                                                                                 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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39; Conserv
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100.0%; Pred. No. 0.0
rative 0; Mismatches
                                                                                                                                              US/09/291,238
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TELECOMMUNICATION INFORMATION:

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RESULT 15
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US-09-291-238-5
 Query Match
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                                                                                                                                                                                                                                                      APPLICATION NUMBER: 09/042,426
FILING DATE: MARCH 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOXIE, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
                                                                                                                                                                                  TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 560 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (919) 541-8689
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/330,760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6239075artis Corporation
                                                                                  HYPOTHETICAL: N
                                                                                                  TOPOLOGICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IMMEDIATE SOURCE:
CLONE: 35S Promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                  IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                322 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 360
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Local Similarity 100.0%;
hes 39; Conservative (
                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: doub
TOPOLOGY: linear
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TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 564 |
CITY: Summit
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Score 39;
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DB 4;
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Length 560;
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Search completed: April 14, 2003, 10:28:30 Job time: 47.5155 secs
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                                                                          322 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 360
                                                                                                           258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296
                                                                                                                                               Pred. No. 0.0081;
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Maximum Match 100%
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Maximum DB seq length: 2000000000
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Perfect score:
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393
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Gapop 10.0 , Gapext 1.0
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:/cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.Seq:*
/cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.Seq:*
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/cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.Seq:*
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                                                                                                                                                                                                                                                                               /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/PC7_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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392	12241	8340	8340	532	515	398	301	371	604	348	541	317	472	600	392	462	393	Length
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US-09-963-803-21	US-10-033-190-5	US-09-874-926-4	US-09-847-057-4	US-09-765-555-1	US-09-963-803-2	US-09-963-803-6	US-09-963-803-7	US-09-963-803-5	US-09-963-803-23	US-09-963-803-4	US-09-963-803-24	US-09-963-803-3	US-09-963-803-25	US-09-963-803-22	US-09-963-803-21	US-09-963-803-20	US-09-963-803-19	ID
Sequence 1, Appli Sequence 21, Appl	1,5	Sequence 4, Appli	,-	Sequence 1, Appli	2	σ,	Sequence 7, Appli	5	23,	4,	_	3, 2	Sequence 25, Appl	Sequence 22, Appl	Sequence 21, Appl	Sequence 20, Appl	Sequence 19, Appl	Description

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US-09-990-659A-15 US-09-810-861B-3	US-09-758-987-1	US-10-162-214-9	US-10-138-221-9	US-09-943-692-33	US-09-951-470-3	US-09-870-375-11	US-09-870-375-6	US-09-870-375-15	US-09-870-375-10	US-09-870-375-9	US-09-870-375-8	US-09-870-375-33	US-09-963-803-13	US-09-870-375-28	US-09-963-803-8	US-09-870-375-5	US-09-870-375-7	US-09-870-375-36	US-09-870-375-35	US-09-870-375-34	US-09-963-803-12	US-09-963-803-9	US-09-963-803-10	TT. CO. CO. 200
Sequence 15, Appl Sequence 3, Appli	Sequence 1, Appli	Sequence 9, Appli	Sequence 9, Appli	Sequence 33, Appl	Sequence 3, Appli	Sequence 11, Appl	Sequence 6, Appli	Sequence 15, Appl	Sequence 10, Appl	Sequence 9, Appli	Sequence 8, Appli	Sequence 33, Appl	Sequence 13, Appl	Sequence 28, Appl	Sequence 8, Appli	Sequence 5, Appli	Sequence 7, Appli	Sequence 36, Appl	Sequence 35, Appl	34	12,	Sequence 9, Appli	Sequence 10, Appl	reduction Tal White

ALIGNMENTS

US-09-963-803-19

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
GENERAL INFORMATION:
ITILE OF INVENTION: Chimeric expression promoters originating from commelina yell
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
FULE REFERENCE: 184332042
FULE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
NORTHER OF SEQ ID NOS: 39
                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 393; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1 SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: promoter LOCATION: (1)..(393)
                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 393
121 AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
                                             13
                                                                                                                           GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACCTAAGCACTGACGAC 120
                                           GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC 120
                                                                                                                                                                                                                 100.0%; Score 393; DB 9; 1 ilarity 100.0%; Pred. No. 2.1e-109; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                          Length 393;
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 199-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 20
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NAME/KEY: promoter
LOCATION: (1)..(462)
OTHER INFORMATION:
S-09-963-803-20
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Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
APPLICANT: Museric expression promoters originating from commelina yellow
TITLE OF INVENTION: Virus and Cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: promoter MPr1163
                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 462
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                                                               181
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           GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
                                                                         AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTT------
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                                                                                                               AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
                                                                                                                           AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
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85.1%;
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Pred. No. 1.9e-85;
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT 1B00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 21
LENGTH: 392
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APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellc
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
FILE REFERENCE: 184332042
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: promoter LOCATION: (1)..(392) OTHER INFORMATION:
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ACGTTACCCGGTATGCCGGTTCCCAAGCTTTATTTCCTTATTTAAGCACTTGTGTAGTAG
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US-09-963-803-25

Sequence 25, Application US/09963803

Publication No. US20030028922A1

GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

TITLE OF INVENTION: Chimeric expression promoters originating

TITLE OF INVENTION: wirus and cassava vein mosaic virus

FILE REFERENCE: 184332042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT FILING DATE: 2001-09-26

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR FILING DATE: 1999-03-29
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Sequence 22, Application US/09963803

Publication No. US20030028922A1

GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

TITLE OF INVENTION: Chimeric expression promoters originating

TITLE OF INVENTION: virus and cassava vein mosaic virus

FILE REFERENCE: 18432042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT FILING DATE: 2001-09-26

PRIOR APPLICATION NUMBER: FR 99/03925
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LOCATION: (1)..(600)
OTHER INFORMATION:
US-09-963-803-22
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LENGTH: 600
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NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
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PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
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Pred. No. 1.5e-82;
0; Mismatches 19
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                                                                                                                                                  SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Matches 310;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                              PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                             APPLICANT: Meristem Therapuetics TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello TITLE OF INVENTION: virus and cassava vein mosaic virus FILE REFERENCE: 184332042
                                                                                                                                                                SOFTWARE: PatentIn version 3.1
                                                                                                                                                                               NUMBER OF SEQ ID NOS: 39
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                                                                                                                 LENGTH: 31
TYPE: DNA
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: promoter LOCATION: (1)..(472) OTHER INFORMATION:
                NAME/KEY: promoter LOCATION: (1)..(31
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                                                FEATURE:
                                                                                                  ORGANISM: Artificial Sequence
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                                                                    OTHER INFORMATION: Promoter Mpr1116
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98.1%;
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PRIOR APPLICATION NUMBER: FR 99/3925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT 1800/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEO ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 24
LENGTH: 541
                                                                                                                            ; NAME/KEY: promoter
; LOCATION: (1)..(541)
; OTHER INFORMATION:
US-09-963-803-24
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Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: Virus and cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 299; Conserv
                                                                               Matches
                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 184332042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                         OTHER INFORMATION: FEATURE:
              138
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                                                        CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
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 Conservative
                                                                                 Conservative
                                                                                                                                                                                   promoter MPr1168
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                                                                                           54.0%;
91.5%;
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79.9%;
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Pred. No. 2.5e-55;
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                                                                              Score 212.4;
Pred. No. 1.3e
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tches 21;
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; NAME/KEY: promoter
; LOCATION: (1)..(348)
; OTHER INFORMATION:
US-09-963-803-4
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                                  US-09-963-803-23
Sequence 23. Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 4
LENGTH: 348
TYPE: DNA
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                                                                                                                                                                                                                                                                                                             Best
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APPLICANT: Meristem Therapuetics TITLE OF INVENTION: Chimeric expuTITLE OF INVENTION: virus and co
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                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION:
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                                                                                                               GGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
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99.1%;
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Pred No. 4.5e-51;
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            promoters originating
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              from commelina yello
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: Vitus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 2000-10-05
NUMBER: OF SEQ ID NOS: 39
NUMBER: OF SEQ ID NOS: 39
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TAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC 199
                                                                                   TGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGA 139
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PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR FILLING DATE: 1999-03-29

PRIOR APPLICATION NUMBER: PCT IB00/00370

PRIOR FILLING DATE: 2000-10-05

NUMBER OF SEQ ID NOS: 39

SOFTWARE: PatentIn version 3.1

SEQ ID NO 23

LENGTH: 604

TYPE: DNA

ORGANISM: Artificial Sequence
                                                                                                           US-09-963-803-5
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; LOCATION: (1)..(604)
; OTHER INFORMATION:
US-09-963-803-23
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APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
TITLE OF INVENTION: Virus and cassava vein mosaic virus
TITLE OF INVENTION: Virus and cassava vein mosaic virus
TITLE OF INVENTION: Virus and cassava vein mosaic virus
TITLE OF INVENTION: Virus and cassava vein mosaic virus
TITLE OF INVENTION: Virus and cassava vein mosaic virus
TITLE OF INVENTION: VIRUS and cassava vein mosaic virus
TITLE OF INVENTION: VIRUS and cassava vein mosaic virus
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TITLE OF INVENTION: VIRUS and cassava vein mosaic
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Best Local Similarity
Matches 221; Conserv
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                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 5
LENGTH: 371
   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 184332042
                                                                                                                                 NAME/KEY: promoter LOCATION: (1)..(371) OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                        LENGTH: 371
TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: promoter MPr1167
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                                                                                                                                                                                                                                                                         OTHER INFORMATION: promoter MPr1146
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGT 304
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
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nilarity 97.4%;
Conservative
39.4%;
75.9%;
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Pred. No. 1.1e-49;
); Mismatches 3;
Score 154.8; DB 9; Pred. No. 3.1e-37;
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                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 7, Application US/09963803 Publication No. US20030028922A1
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: ECT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Chimeric expression pron
TITLE OF INVENTION: virus and cassava vein
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Meristem Therapuetics
                                                                                                                                                                                                                                                                                                                                              LENGTH: 30
TYPE: DNA
                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                     FEATURE:
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182
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GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
                GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACAACCTAGAGGATCC 371
                                                       GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
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                                                                                                                                                                                    Similarity
                                                                                                                                                                        Conservative
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98.1%;
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                                                                                                                                                                      Score 140.8; DB 9;
Pred. No. 4.9e-33;
0; Mismatches 2;
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; OTHER INFORMATION: promoter MPr1147; FEATURE: NAME/KEY: promoter; LOCATION: (1)..(398); OTHER INFORMATION: US-09-963-803-6
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US-09-963-803-2
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                                                                                                                                                                                                                                                                                                                  Sequence 2, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT TAPPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
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NUMBER OF SEC ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEC ID NO 6
LENGTH: 398
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Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 153; Conserv
                                                                                                                                                                                 SEQ ID NO 2
LENGTH: 515
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05
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                                                                   OTHER INFORMATION: Promoter from the intergenic region of Cassava Vein Mosaic virus OTHER INFORMATION: of 515 bp in length EMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                   FEATURE:
                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE
LOCATION:
                     NAME/KEY:
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(1)...(515)
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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Methods and composition:
TITLE OF INVENTION: expression in plants
FILE REFERENCE: 27801-20014.40
CURRENT APPLICATION NUMBER: US/09/765,555
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: US 09/620,897
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 2000-01-21
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                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 532
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Publication No. US20030037355A1
GENERAL INFORMATION:
APPLICANT: The Scripps Research Institute
                                                                                                            Sequence 4, Application US/09847057 Patent No. US20020004943A1
                                                                                                                                                                                                                                                                                                                                                                                                   Matches 126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 126;
APPLICANT: AGRINOMICS, LLC.
TITLE OF INVENTION: IDENTIFICATION AND CHARACTERIZATION OF A PAGODA PHENOTYPE IN FILE REFERENCE: PAGODA
CURRENT APPLICATION NUMBER: US/09/847,057
CURRENT FILING DATE: 2001-05-01
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                    Best
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                      153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93 GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT 152
                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                      ATCACAAAGGAATCT 365
                                                                                                                                                                                                                                         ATGCATTTGTAATTT 227
                                                                                                                                                                                                                                                                                                                                       GAATCTTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT
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93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 120.6; DB y;
Pred. No. 8.3e-27;
Pred. No. 8.3e-27;
                                                                                                                                                                                                                                                                                                                                                                                                  Score 120.6; DB 9; Pred. No. 8.4e-27; 0; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    compositions to modulate
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                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 532;
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; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 4
; LENGTH: 8340
; TYPE: DNA
; TYPE: DNA
; ORGANISM: Binary vector pAGI4002
US-09-847-057-4

Query Match
Best Local Similarity 93.3%; Pred. No. 3.1e-26;
Matches 126; Conservative 0; Mismatches 9; Indels 0; Gaps 0;
Matches 126; Conservative Adamanaranagananaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagaranagarana
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Maximum DB seq length: 2000000000
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Perfect score:
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Gapop 10.0 , Gapext 1.0
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            cgn2_6/ptodata/1/pna/US06_COMB.seq:*
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cgn2_6/ptodata/1/pna/PCTUS_COMB.seq
/cgn2_6/ptodata/1/pna/US101B_COMB.seg
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6232.082 Million cell updates/sec
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44: /cgn2_6/ptcdata/1/pna/US6001_COMB.seq:
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47: /cgn2_6/ptcdata/1/pna/US6003_COMB.seq:
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53: /cgn2_6/ptcdata/1/pna/US60101_COMB.seq:
54: /cgn2_6/ptcdata/1/pna/US6011_COMB.seq:
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70: /cgn2_6/ptcdata/1/pna/US6022_COMB.seq:
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77: /cgn2_6/ptcdata/1/pna/US6023_COMB.seq:
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79: /cgn2_6/ptcdata/1/pna/US6033_COMB.seq:
79: /cgn2_6/ptcdata/1/pn
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

20 21	19	18	17	16	15	14	13	12	11	10	9	8	7	6	տ	4	ω	2	ᆫ	No.	Result
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30.7 30.7	30.7	30.7	30.7	30.7	30.7	30.7	30.7	35.8	35.8	39.4	49.9	51.0	54.0	54.6	72.6	77.5	78.9	79.9	100.0	Match Length	Query
411 515	411	411	411	392	392	392	392	398	301	371	604	348	541	317	472	600	392	462	393		
16 25	16	_	_	16	16	1	_	36	36	36	36	36	36	36	36	36	36	36	36	DB -	
US-09-202-838A-4 US-09-641-466-1	US-09-202-838-4	PCT-US97-10376A-4	PCT-US97-10376-4	US-09-202-838A-1	US-09-202-838-1	PCT-US97-10376A-1	PCT-US97-10376-1	US-09-963-803-6	us-09-963-803-7	us-09-963-803-5	US-09-963-803-23	US-09-963-803-4	US-09-963-803-24	US-09-963-803-3	US-09-963-803-25	US-09-963-803-22	US-09-963-803-21	US-09-963-803-20	US-09-963-803-19	ID	
Sequence 4, Appli Sequence 1, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 6, Appli	Sequence 7, Appli	Sequence 5, Appli	Sequence 23, Appl	Sequence 4, Appli	Sequence 24, Appl	Sequence 3, Appli	Sequence 25, Appl	-	Sequence 21, Appl	-	Sequence 19, Appl	Description	

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Sequence 19, Application US/09963803

GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

ITILE OF INVENTION: Chimeric expression promoters originating from the respective of the promoter of the promote
                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: promoter MPr1162
FEATURE:
NAME/KEY: promoter
LOCATION: (1)..(393)
OTHER INFORMATION:
US-09-963-803-19
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Best Local Similarity
Matches 393; Conser
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                 AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
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US-09-202-838A-3
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US-09-75-555-1
US-09-641-466-2
US-09-673-274A-20
US-09-673-274A-20
US-09-673-274A-20
US-09-874-926-4
US-09-874-926-4
US-09-874-926-4
US-09-874-926-4
US-09-874-926-4
US-09-874-926-888-5
US-09-987-10376A-5
PCT-US97-10376A-9
PCT-US97-10376A-9
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PCT-US97-10376-2
PCT-US97-10376A-2
6 US-09-202-838-2
6 US-09-202-838-3
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Pred. No. 2.6e-103;
); Mismatches 0;
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; LOCATION: (1)..(46:
; OTHER INFORMATION:
US-09-963-803-20
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GENERAL INFORMATION:
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Best Local Similarity
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TYPE: DNA
ORGANISM: Artificial Sequence
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PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 20
LENGTH: 462
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
                                                                                                                                                                                                                                                                                            OTHER INFORMATION: promoter MPr1163
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                                    CTTAGAAAACCAACAACAACCTAGAGGATCC
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                                                                                                                Score 314; DB 36;
Pred. No. 2.5e-80;
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US-09-963-803-21
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
FILE REFERENCE: 184332042
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SEQ ID NO 21
LENGTH: 392
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: promoter LOCATION: (1)..(392) OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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                                                           CTTAGAAAACCAACAACAACAAC 382
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337; Conserv
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88.2%;
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Pred. No. 3.4e-79;
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  SOFTWARE: Pat
SEQ ID NO 25
LENGTH: 472
TYPE: DNA
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Best Local (
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; NAME/KEY: promoter
; LOCATION: (1)..(600)
; OTHER INFORMATION:
US-09-963-803-22
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                                                                                                                                                                                                                                                                                Sequence 25, Application US/09963803 GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters origina TITLE OF INVENTION: virus and cassava vein mosaic virus FILE REFERENCE: 184332042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT FILING DATE: 2001-09-26

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR FILING DATE: 1999-03-29

PRIOR APPLICATION NUMBER: PCT IB00/00370

PRIOR FILING DATE: 2000-10-05

NUMBER OF SEQ ID NOS: 39

SOFTWARE: Patentin version 3.1
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-01-095
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
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TYPE: DNA
ORGANISM: Artificial Sequence
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316; Conserv
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Pred. No. 1.5e-77
0; Mismatches 1
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PATENTIN VETSION 3.1
SEQ ID NO 3
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                                                                                                         US-09-963-803-3
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
                                                                              Query Match
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Best Local Similarity
Matches 310; Conserv
                                                      Matches
                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow TITLE OF INVENTION: virus and cassava veln mosaic virus FILE REFERENCE: 184332042
                                                                                                                   NAME/KEY: promoter LOCATION: (1)..(317) OTHER INFORMATION:
                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                          FEATURE:
                                                                                                                                                                     OTHER INFORMATION: Promoter MPr1116
                                                                                                                                                                                  FEATURE:
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Local Similarity
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                                                     Conservative
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Pred. No. 5.8e-72;
                                                               Score 214.4; DB 3 Pred. No. 1.9e-51;
                                                     Mismatches
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; LOCATION: (1)..(541)
; OTHER INFORMATION:
US-09-963-803-24
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US-09-963-803-24
; Sequence 24, Application US/09963803
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LENGTH: 54
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Best Local Similarity
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APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                      279
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             258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACCCGGTATGCC 317
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                                                   AACCTAGAGGATCC 393
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 TTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACCCGGTATGCCGG
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91.5%;
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Pred. No. 8.7e-51;
0; Mismatches 21;
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FEATURE:

NAME/KEY: promoter

LOCATION: (1)..(348)

OTHER INFORMATION:
US-09-963-803-4
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
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 NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
                                                                                                                                                                                                Sequence 23, Application US/09963803 GENERAL INFORMATION:
                                    APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR PILING DATE: 2000-10-05
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Best Local Similarity
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APPLICANT: Meristem Therapuetics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGATATCAAGATTGATGT
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99.1%;
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; NAME/KEY: promoter ; LOCATION: (1)..(371) ; OTHER INFORMATION: US-09-963-803-5
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US-09-963-803-5
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NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5, Application US/09963803 GENERAL INFORMATION:
                                                                                                                                  Query Match
Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
                                                                                                                                                                                                                                                                                                                                  LENGTH: 371
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                              FEATURE
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OTHER INFORMATION: promoter MPrll46
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138 GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197
                                       130 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 189
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                                                         78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
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75.98;
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97.48;
                                                                                                               Score 154.8; DB 36;
Pred. No. 4.2e-34;
0; Mismatches 2;
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Pred. No. 4.6e-46;
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249 257

190 GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG

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PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
                                                                                                                                                                                            RESULT 12
US-09-963-803-6
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                                                                                  Sequence 6, Application US/09963803
GENERAL INFORMATION:
APPLICANT: METISLE Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
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APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
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Best Local Similarity
Matches 153; Conserv
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CURRENT APPLICATION NUMBER: US/09/963,803 CURRENT FILING DATE: 2001-09-26 PRIOR APPLICATION NUMBER: FR 99/03925 PRIOR FILING DATE: 1999-03-29 PRIOR APPLICATION NUMBER: PCT IB00/00370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                   78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                              GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 217
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Pred. No. 4.5e-30;
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MOLECULE TYPE: DN
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                                                  153 TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT
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CLASSIFICATION:
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: NAME/KEY: promoter
: LOCATION: (1)..(398)
: OTHER INFORMATION:
US-09-963-803-6
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NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 6
LENGTH: 398
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 20-JUN-1 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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DEDNESS: single
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93.3%;
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98.1%;
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Score 120.6; DB 1,
Pred. No. 3.7e-24;
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                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (619) 784-93 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 20-JUN-1996
ATTOREY/AGENT INFORMATION:
NAME: FITLING, Thomas
REGISTRATION NUMBER: 34,163
REGERENCE/DOCKET NUMBER: 504
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 784-2937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US97/10376A FILING DATE: 20-JUN-1997 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 392 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: CASSAVA VEIN MOSAIC VIRUS PROMOTERS TITLE OF INVENTION: USES THEREOF NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Verdaguer, Bertrand
APPLICANT: de Kochko, Alexandre
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                                                                                                                                                                                                      30.7%;
Local Similarity 93.3%;
ses 126; Conservation
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: La Jolla
STATE: California
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TOPOLOGY: lin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                 ATGCATTTGTAATTT 227
                                                                                                   TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT 212
ATCACAAAGGAATCT 277
                                                                    TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT 262
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Pred. No. 3.7e-24;
                                                                                                                                                                                                            Mismatches
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RESULT 15

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Search completed: April 14, 2003, 12:06:29 Job time: 1588.51 secs
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; TYPE: DNA
; ORGANISM: Cassava vein mosaic virus
US-09-202-838-1
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GENERAL INFORMATION:
APPLICANT: Verdaguer, Bertrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/202,838
CURRENT FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/US97/10376
PRIOR FILING DATE: 1997-06-20
PRIOR APPLICATION NUMBER: 60/020,129
PRIOR FILING DATE: 1996-06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: MYC0011S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Fauquet, Claude
TITLE OF INVENTION: CASSAVA VEIN MOSAIC VIRUS PROMOTERS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                            263
                                                                                                                                                                                                                             143 GAATCTTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT
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                                                                            ATCACAAAGGAATCT
                                                                                                             ATGCATTTGTAATTT 227
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de Kochko, Alexandre
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93.3%;
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Pred. No. 3.7e-24;
0; Mismatches 9;
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Match
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      Length
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US-09-765-558-1
US-09-673-274B-19
US-09-673-274B-20
US-09-673-274B-20
US-09-673-274B-20
US-09-949-016-13294
US-10-380-935-30
US-10-380-935-37
US-10-391-414-10
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US-10-130-150-13
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US-10-160-764-40
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Sequence 20, Appl
Sequence 30, Appl
Sequence 37, Appl
Sequence 37, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 117, Appl
Sequence 1, Appl
Sequence 40, Appl
Sequence 40, Appl
Sequence 52, Appl
Sequence 53, Appl
Sequence 54, Appl
Sequence 56, Appl
Sequence 57, Appl
Sequence 58, Appl
Sequence 59, Appl
Sequence 59, Appl
Sequence 51, Appl
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113,	Sequence 893, App	7 ,	, 0	Sequence 17, Appl	Sequence 57, Appl	•		Sequence 7, Appli		Sequence 31, Appl	•	•	•	Sequence 38, Appl	•	Sequence 11, Appl	Sequence 6, Appli	Sequence 22, Appl	Sequence 9, Appli		Sequence 13, Appl

Title: Perfect score:

Sequence:

Scoring table:

ALIGNMENTS

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APPLICANT: Guan, Xueni
APPLICANT: Dalmia, Bipin
ITITLE OF INVENTION: Methods and compositions to modulate
ITITLE OF INVENTION: expression in plants
FILE REFERENCE: 27801-20014.20
CURRENT APPLICATION NUMBER: US/09/765,555A
CURRENT FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 09/620,897
PRIOR APPLICATION NUMBER: US 60/177,468
PRIOR FILING DATE: 2000-07-21
PRIOR FILING DATE: 2000-01-21
VALUE OF SEQ ID NOS: 75
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
FENGTH: 532
                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Promoter CsVMV US-09-765-555A-1
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US-09-765-555A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 1, Application US/09765555A
; GENERAL INFORMATION:
                                                                                                                                                                                                                                        Query Match 30.7%;
Best Local Similarity 93.3%;
Matches 126; Conservative
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APPLICANT: Barbas, Carlos
APPLICANT: Stege, Justin
APPLICANT: Guan, Xueni
                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                         351 ATCACAAAGGAATCT 365
                                     ATGCATTTGTAATTT 227
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Pred. No. 1.2e-25;
0; Mismatches 9;
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Result

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Score

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Sequence 19, Application US/09673274B GENERAL INFORMATION:

09-673-274B-19

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APPLICANT: Lamberty, Mireille
APPLICANT: Bulet, Phillipe
APPLICANT: Bulet, Phillipe
APPLICANT: Bulet, Phillipe
APPLICANT: Hoffman, Jules
TITLE OF INVENTION: GENE CODING FOR HELIOMICINE, ANIFILE REFERENCE: A33595-PCT'USA (072667.0166)
CURRENT APPLICATION NUMBER: US/09/673,274B
CURRENT FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: PCT/FR99/00843
PRIOR APPLICATION NUMBER: FR 98 04933
PRIOR FILING DATE: 1998-04-15
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Best Local
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 19
                                     SOFTWARE: FO
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APPLICANT: Brokhart, Gary
APPLICANT: Brokhart, Gary
APPLICANT: Hoffman, Jules
TITLE OF INVENTION: GENE CODING FOR HELIOMICINE, AND USE THEREOF
FILE REFERENCE: A33595-PCT-USA (0726667.0166)
CURRENT APPLICATION NUMBER: US/09/673,274B
CURRENT FILING DATE: 1999-04-12
PRIOR FILING DATE: 1999-04-12
PRIOR FILING DATE: 1999-04-12
PRIOR PRIOR FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: FR 98 04933
PRIOR FILING DATE: 1998-04-15
PRIOR FILING DATE: 1998-04-15
                                                                                  NUMBER OF SEQ ID NOS:
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ORGANISM: Artificial Sequence
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local to 126;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATCACAAAGGAATCT 356
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                      1036
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26; Conservative
                                                              FastSEQ for Windows Version 4.0
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Pred. No. 1.4e
0; Mismatches
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(250715)
OTHER INFORMATION: n = A
US-09-949-016-13294
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                                                                                                             Db 146653
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                                                                                                                                                                                      Query Match
Best Local
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SEQ ID NO 13294
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Best Local Similarity
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR ETLING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/241,755 PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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                                                                             179 TAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGACTA 238
                                                                                                                                          119 ACAACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATG
239 GTGATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGA 289
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                                                                                                                                                                                    Local
                                                                                                            ATGCATTTGTAATTT 227
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93.3%;
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Pred. No. 1.
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OF DETECTION AND USES THEREOF
                                                                                                                                                                                                    Length 250715;
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; OTHER INFORMATION: Enhancer US-10-380-935-37
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US-10-380-935-30
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                                                                         Query Match
Best Local Similarity
Matches 39; Conserv
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PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: GB0027693.1
PRIOR EILING DATE: 2000-11-13
PRIOR APPLICATION NUMBER: GB0023910.3
PRIOR APPLICATION NUMBER: GB0023910.3
PRIOR FILING DATE: 2000-09-29
                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: GB0023911.1
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: GB0027693.1
PRIOR FILING DATE: 2000-11-13
PRIOR PPLICATION NUMBER: GB0023910.3
PRIOR APPLICATION NUMBER: GB0023910.3
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/380,935
CURRENT FILING DATE: 2003-03-19
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APPLICANT: HAWKES, Timothy, Robert
APPLICANT: ANDREWS, Christopher, John
TITLE OF INVENTION: HERBICIDE RESISTANT PLANTS
FILE REFERENCE: PPD50594/WO
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CURRENT FILING DATE: 2003-03-19
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APPLICANT: ANDREWS, Christopher, John
TITLE OF INVENTION: HERBICIDE RESISTANT PLANTS
FILE REFERENCE: PPD50594/WO
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                                                                                                                                                                                                                  LENGTH: 522
TYPE: DNA
ORGANISM: Figwort Mosaic Virus
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    459 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 497
                                          258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296
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                                                                                 Conservative
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                                                                                                  9.9%; Score 39; DB 8; 100.0%; Pred. No. 0.2;
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                                                                                 Mismatches
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0.17;
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US-10-302-267-1; Sequence 1, Application US/10302267; GENERAL INFORMATION:
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                                                             RESULT
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; ORGANISM: Cauliflower mosaic virus
US-10-391-414-10
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US-10-391-414-10
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APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES IN PLANTS
FILE REFERENCE: MBI-0051 P
CURRENT APPLICATION NUMBER: US/60/434,166
CURRENT FILING DATE: 2002-12-17
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APPLICANT: ITO, YUSUKE
TITLE OF INVENTION: Stress Induced Promoter Derived From Rice
FILE REFERENCE: 382.1041
CURRENT APPLICATION NUMBER: US/10/391,414
CURRENT FILING DATE: 2003-03-18
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PRIOR FILING DATE: 2002-12-26
NUMBER OF SEQ ID NOS: 11
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APPLICANT: KATSURA, Koji
                                                                                                                                                                                                                                                                                                                            LENGTH: 1683
TYPE: DNA
ORGANISM: Arabidopsis thaliana
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APPLICANT: Creelman, Robert A.
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                                                                                                                                                         258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296
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                                                                                                                                                                                                                  Local
                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jiang, Cai-Zhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Haake, Volker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ratcliffe, Oliver
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100.0%; Pr
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; Pred. No.
                                                                                                                                                                                                                  Score 39; DB 11;
Pred. No. 0.27;
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0.22;
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Keddie, James

Fromm, Michael

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RESULT 10
US-10-130-150-13
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                                                                                                                                                                     Sequence 13, Application US/10130150
GENERAL INFORMATION:
APPLICANT: Shoseyov, Oded
TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THEREOF
FILE REFERENCE: 02/23831
                                                    SOFTWARE: PatentIn version 3.1
SEQ ID NO 13
LENGTH: 3212
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Best Local :
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SEQ ID NO 1
LENGTH: 2361
                                                                                                            CURRENT APPLICATION NUMBER: US/10/130,150
CURRENT FILING DATE: 2000-11-15
NUMBER OF SEQ ID NOS: 24
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/120,880
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: 60/121,037
PRIOR FILING DATE: 1999-02-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pilgrim, Marsha
APPLICANT: CreeLman, Robert
TITLE OF INVENTION: PLANT GENE SEQUENCES II
FILE REFERENCE: MBI-0007
CURRENT APPLICATION NUMBER: US/10/302,267
CURRENT FILING DATE: 2002-11-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1999-11-01
NUMBER OF SEQ ID NOS: 218
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ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                  258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296
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OR FILING DATE: 1999-03-11
OR APPLICATION NUMBER: 60/129,450
OR FILING DATE: 1999-04-15
OR APPLICATION NUMBER: 60/135,134
OR FILING DATE: 1999-05-20
OR APPLICATION NUMBER: 60/144,153
OR APPLICATION NUMBER: 60/144,153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/161,143 FILING DATE: 1999-10-22
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Reuber, Lynne
Zhang, James
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Riechmann, Jose Luis
Adam, Luc
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Jiang, Cai-Zhong
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100.0%; Pred. No.
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0.3;
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              RESULT 13
US-10-160-764-40
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Sequence

40, Application US/10160764

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; OTHER INFORMATION: Sequence of recombinant construct expressing BGL1 protein fuse ; OTHER INFORMATION: and ER retaining signal peptides US-10-130-150-18
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                                                                                                                                 US-10-130-150-15
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US-10-130-150-15
                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/130,150
CURRENT FILING DATE: 2000-11-15
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.1
SEQ ID NO 15
LENGTH: 3329
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1
SEQ ID NO 18
LENCTH: 3288
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                              Sequence 15, Application US/10130150 GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity 100.0%;
                                                               Matches
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TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THER
EILE REFERENCE: 02/23831
CURRENT APPLICATION NUMBER: US/10/130,150
CURRENT FILING DATE: 2000-11-15
                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 24
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                                                                                                                                               OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial sequence
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197
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es 39; Conserv
GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 235
                                                               39;
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                                                                                                                                           Sequence of recombinant construct expressing BGL1 protein fuse signal peptide
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0; Mismatches
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Pred. No.
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0.33;
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US-10-160-764-4
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OTHER INFORMATION: Description of Artificial Sequence: Plasmid
OTHER INFORMATION: pBI121-35S-AtETA
US-10-160-764-40
                                                                                                                                                                US-10-160-764-4
                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR FILING DATE: 2001-10-22
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PATENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Huang, Ya
APPLICANT: Chalifou
APPLICANT: Wang, Ya
                                                                                                                                                                                                                          SEQ ID NO 4
LENGTH: 5250
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 40
LENGTH: 5247
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                  Query Match
Best Local Similarity
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TITLE OF INVENTION: Compositions and Methods of Increasing Stress Tolerance
TITLE OF INVENTION: in plants
FILE REFERENCE: 22542-008
CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR PILING DATE: 2001-10-22
NUMBER OF SEQ ID NOS: 90
NUMBER OF SEQ ID NOS: 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 22542-008
CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
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APPLICANT: Gilley, Angela Patricia
TITLE OF INVENTION: Compositions and Methods of Increasing Stress Tolerance
TITLE OF INVENTION: in Plants
                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Plasmid OTHER INFORMATION: pBI121-355-anti-AtFTA sequence
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3209 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 3247
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                   258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296
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Local Similarity 100.0%; Pred. No.
les 39; Conservative 0; Mismatc
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ilarity 100.0%;
Conservative (
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                                                                                                Score 39; DB 9;
Pred. No. 0.37;
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0.37;
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                                                                                                                     Length 5250;
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; OTHER INFORMATION: Description of Artificial Sequence: Plasmid ; OTHER INFORMATION: pBI121-35S-Anti-AtFTB US-10-160-764-48
Search completed: April 14, 2003, 12:27:28 Job time: 407.159 secs
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US-10-160-764-48
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TITLE OF INVENTION: Compositions and Methods
TITLE OF INVENTION: Compositions and Methods
TITLE REFERENCE: 22542-008
CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR FILING DATE: 2001-10-22
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                       Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 48, Application US/10160764 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                               SEQ ID NO 48
LENGTH: 5511
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Huang, Yafan APPLICANT: Chalifoux, Maryse
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                                                                            3209 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 3247
                                                                                                 258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296
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100.0%; Pred. No.
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0.38;
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Title: Perfect score: Sequence:

US-09-963-803-19 393 1 aagcttgcatgcctgcagca.....cacaacaacctagaggatcc 393 April 14, 2003, 06:53:52; Search time 913.489 Seconds (without alignments) 6967.603 Million cell updates/sec

Run on:

OM nucleic - nucleic search, using sw model

GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

Scoring table:

IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

em_estba:*
em_esthum:*
em_estin:*

em_estmu:*
em_estov:*
em_estpl:*
em_estro:*
em_htc:*

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Total number of hits satisfying chosen parameters:

32308132

16154066 seqs, 8097743376 residues

Searched:

0 00 0	0000000	0 00 0	0 0
00000000000000000000000000000000000000	33333333333333333333333333333333333333	18 19 20 20 21 22 23 24 25 26 27 28	7 9 10 11 12 12 13 14 15
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0		U U U U U U U U U U U U U U U U U U U	
274 279 281 281 284 284 296	256 258 261 268 269 271 272	215 219 220 220 230 237 244 248 251	112 142 153 158 158 175 177 190
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BH802495 BH748475 BH750170 BH802443 BH213307 BH213307 BH747007 BH747007 BH610310 BH213532	BH748499 BH802470 BH802493 BH611919 BH211646 BH802489 BH799178 BH802428	BH753813 BH747744 BH747744 BH747438 BH802415 BH746474 BH799173 BH802463 BH802463 BH746375 BH746375	BH751161 BH749349 BH619283 BH747013 BH747013 BH748291 BH747357 BH746616 BH802465 BH752801 BH752801 BH74829
BH802495 BH748475 BH750170 BH802443 BH213307 BH746665 BH747007 BH610310 BH213532	BH748499 BH802470 BH802493 BH611919 BH211646 BH802489 BH799178 BH802428	BH753813 BH747744 BH747438 BH802415 BH746474 BH799173 BH802463 BH802463 BH76375 BH254798 BH750171	BH751161 BH749349 BH619283 BH7470133 BH748291 BH747357 BH746616 BH8024655 BH752801 BH752801
1008026H0 SALK_0460 SALK_0371 1008026D0 SALK_0090 SALK_0097 SALK_0078 SALK_0078 SALK_0085 SALK_0093	SALK_0460 11008026F0 11008026H0 SALK_0318 SALK_0064 11008026G0 11008025C0	SALK 0296 SALK 0337 SALK 0174 1008026B0 SALK 0428 1008025B1 1008025E0 SALK 0402 SALK 0402 SALK 04037 SALK 0371	SALK_0494 SALK_0477 SALK_04077 SALK_04080 SALK_0451 SALK_0165 SALK_0452 1008026E1 SALK_0193 SALK_0450

ALIGNMENTS

RESULT 1	
BH746858	
LOCUS	BH746858 224 bp DNA linear GSS 27-FEB-2002
DEFINITION	SALK_003694.51.40.x Arabidopsis thaliana TDNA insertion lines
	Arabidopsis thaliana genomic clone SALK_003694.51.40.x, DNA
	sequence.
ACCESSION	BH746858
VERSION	BH746858.1 GI:18959973
KEYWORDS	GSS.
SOURCE	thale cress.
ORGANISM	Arabidopsis thaliana
	Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
	Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
	Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
REFERENCE	1 (bases 1 to 224)
AUTHORS	Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., Gadrinab
	,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P.
	, Zimmerman, J. and Ecker, J.R.
TITLE	A Sequence-Indexed Library of Insertion Mutations in the
	Arabidopsis Genome
JOURNAL	Unpublished (2001)
COMMENT	Contact: Joseph R. Ecker
	Salk Institute Genomic Analysis Laboratory (SIGnAL)
	The Salk Institute for Biological Studies
	10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
	Tel: 858 453 4100 x1752
	Fax: 858 558 6379
	Email: ecker@salk.edu
	This is single pass sequence recovered from the left border of

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

em_gss_mam: *
em_gss_mus: *
em_gss_other: *
em_gss_pro: *
em_gss_rod: *

em_estfun:*

em_estom:*
gb_gss:*
em_gss_hum:*
em_gss_inv:*
em_gss_pin:*
em_gss_vrt:*
em_gss_fun:*

gb_est1: *
gb_est2: *
gb_htc: *
gb_est3: *
gb_est4: *
gb_est5: *

SUMMARIES

c 6	U	4	ω	2	1	Result No.
39.2	40	40.2	41	42.4	42.4	Score
10.0	10.2	10.2	10.4	10.8	10.8	Query Match Length DB ID
480	911	528	222	561	224	Length DB ID
17	17	9		13	17	DB:
вн639017	AZ548969	AL369550	BM161725	BM162517	BH746858	Ħ
вн639017 1008026н0	AZ548969 ENTEM27TF	AL369550 MtBA31G10	BM161725 EST564248	BM162517 EST565040	BH746858 SALK_0036	Description

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VERSION
KEYWORDS
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BM162517
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Best Local Similarity 81.7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Jane Cariton
Parasite Genomics Group
The Institute for Genomic Research
9712 Medical Center Drive, Rockville, MD 20850, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carlton, J.M., Daly, T.M., Long, C.A., Bergman, L.W., Vaidya, A.B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Plasmodium yoelii yoelii
Lukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              For clone info, please contact the Malaria Research and Reference Reagent Resource Center, ATCC http://www.malaria.mr4.org/mr4pages/index.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: carlton@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fraser, C.M. and Carucci, D.J.
Plasmodium yoelii EST project at TIGR
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/dev_stage="Assaval blood stages"
//lab_host="E. coli XL-1 Blue"
//lab_host="E. coli XL-1 Blue"
//note="Vector: pAD-GAL4; At 20-25% parasitemia, blood was collected from BALB/CBVJ mice infected with Py17XL parasites, and leukocytes removed by passage over microcrystalline cellulose collumns. Total RNA was isolated using the guanddinium isothiocyanate method, and mRNA isolated using oligo(dT)-cellulose chromatography. First strand cDNA synthesis was completed using a 50-base primer and reverse transcriptase in the presence of 5-methyl dCTP. After second strand synthesis, uneven termini were treated with Pfu DNA polymerase and EcoRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Arabidopsis thallana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thallana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 68 c 54 g 50 t
                                                                                                                                                                                                                                                                                                                                                                    /organism="Plasmodium yoelii yoelii"
/strain="17XL"
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/clone="SALK_003694.51.40.x"
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/strain="Columbia 0"
                                                                                                                                                                                                                                                                                                /clone_lib="PyBS"
                                                                                                                                                                                                                                                                                                                   /clone="PYCKT64"
                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:73239"
                                                                                                                                                                                                                                                                                                                                                                                                                                     location/Qualifiers
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Pred. No. 0.83;
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PYCKT64 5' end,
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RESULT 3
BM161725
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            69 ATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGA 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Parasite Genomics Group
The Institute for Genomic Research
9712 Medical Center Drive, Rockville,
Tel: 301-530-9319
Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmodium yoelii EST project at Unpublished (2001)
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EST564248 PyBS Plasmodium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium. 1 (bases 1 to 222)
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BM161725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    For clone info, please contact the Malaria Research and Reference Reagent Resource Center, ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Jane Carlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fraser, C.M. and Carucci, D.J
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                                                                                                                                                                                                                                                                                                                                                                                                                         http://www.malaria.mr4.org/mr4pages/index.html
Seq primer: ADF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: carlton@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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/dev_stage="Asexual blood stages"
/lab_host="E. coli XL-1 Blue"
/lab_host="E. coli XL-1 Blue"
/lab_host="E. coli XL-1 Blue"
/note="Vector: pAD-GAL4; At 20-25% parasitemia, blood was collected from BALB/CByJ mice infected with Py17XL parasites, and leukocytes removed by passage over microcrystalline cellulose collumns. Total RNA was isolated using the guanidinium isothiocyanate method, and mRNA isolated using oligo(dT)-cellulose chromatography. First strand cDNA synthesis was completed using a 50-base primer and reverse transcriptase in the presence of 5-methyl dCTP. After second strand synthesis, uneven
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82 g 183 t
                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                              /clone_lib="PyBS"
                                                                                                                                                                                                                                                                                      /clone="PYCKI31"
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                                                                                                                                                                                                                                                                                                                                      /strain-"17XL"
                                                                                                                                                                                                                                                                                                                                                         /organism="Plasmodium yoelii yoelii"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88 TGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGTAAAAAGATAGAGTATATATATAACAACAACAGTCGTGCTGAAAATAACGCAAAATCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                              BP 191 91006 EVRY cedex - France
Email: segref@genoscope.cns.fr. Web : www.genoscope.cns.fr
Contact : Pascal Gamas and Etienne-Pascal Journet, Laboratoire
Biologie Moleculaire des Relations Plantes Microorganismes,
CNRS-INRA, BP 27 31326 Castanet-Tolosan Cedex, France (Email :
Mt-est@toulouse.inra.fr Website :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AL369550 528 bp mF
MtBa31G10R1 MtBA Medicago truncatula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Medicago truncatula ESTs from nitrogen-starved Unpublished (2000)
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Journet, E.P., Crespeau, H., van-Tuinen, D., Gouzy, J., Jaillon, O., Niebel, A., Carreau, V., Chatagnier, O., Kahn, D., Gianinazzi-Pearson
                                                                                                                                                                                                                                                                                                                                                                                                                                 http://sequence.toulouse.inra.fr/Mtruncatula.html).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and Gamas, P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                               /tissue_type="root tips"
/dev_stage="harvested after 3 days of N-starvation"
/dev_stage="harvested after 3 days of N-starvation"
/note="vector: pBluescript pSK; Site_1: EcoR; Site_2:
XhoI; Plants were grown in an aeroponic chamber for 14
days on nitrogen-rich medium followed by 3 days on N-free
medium. RNA was extracted from root tips (1-3 cm). cDNA
was prepared from polyA+ enriched RNA. The CDNA was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          isolated."
1 25 c
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                          directionally ligated into Uni-zapXR vector from Stratagene and packaged using Gigapack Gold packa
                                                                                                                                                                                                                                                                                 /db_xref="taxon:3880"
/clone="MtBA31G10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        adaptors ligated to the blunt ends. The sample was cleaved with XhoI and separated on a Sephacryl S-500 column.
                                                                                                                                                                                                                                                                                                                             /cultivar="Jemalong"
                                                                                                                                                                                                                                                    /clone_lib="MtBA"
                                                                                                                                                                                                                                                                                                                                                        ∕organism="Medicago truncatula"
                                                                                                                                                                                                                                                                                                                                                                                                         ocation/Qualifiers
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Pred. No. 1.9;
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  Gigapack Gold packaging cDNA inserts were
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   254 AAAGACTTTGATGAAGATGGTTGTCATGGAATAGAGTATTATGATCCATTGAAGGCTAAG 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAAAAAAAAAAAACTCGAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACC
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AZ548969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AZ548969 911 bp DNA linear GSS 14-NOV-200 ENTEM27TF Entamoeba histolytica Sheared DNA Entamoeba histolytica
                                                                                                                                                                                                                                       High quality sequence start: High quality sequence stop: 7
                                                                                                                                                                                                                                                                              Seq primer: M13-Forward
Class: shotgun
                                                                                                                                                                                                                                                                                                                                        Email: bjloftus@tigr.org
Clones are derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Department of Eukaryotic Genomics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Brendan J Loftus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
                                                                                                                                                                                                                                                                                                                         DNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Determination of clone end sequences HM1:IMSS sheared DNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Entamoebidae; 1 (bases 1 to 911)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Entamoeba histolytica
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Loftus, B., Van Aken, S. and Fraser, C
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301 838 3543
                                                                                                                                                                                                                                                                                                                                                                                                                    Medical Center Dr.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                /clone_lib="Entamoeba histolytica Sheared DNA"
/note="Vector: pHOS1; Site_l: Bst I; Constructed at The
Institute for Genomic Research (TIGR), Rockville, MD.
Genomic DNA isolated from broth cultures of E. histolytic
using a method described by Clark and Diamond (Clark,
C.G., and Diamond, L.S. (1993) Entamoeba histolytica: a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mass-excised from phage stocks using ExAssit helper phage and propagated in SOLR cells. Clone ordering and sequencing was performed by the Centre National de Sequencage (Genoscope, Evry, France)."
                                                                                                                                        /strain="HM1:IMSS"
/db_xref="taxon:5759"
                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                               /organism="Entamoeba histolytica"
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47.1%;
and Diamond, L.S. (1993) En
d for isolate identification.
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Pred. No. 3
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: walbot@stanford.edu
Possible ligation site so sequence
sequence submitted separately.
                                                                                                                                                                                                                                                                                                                                                                                       Stanford University
855 California Ave, Palo
                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2001 Contact: Walbot V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Walbot, V.
Maize genomic sequences found using engineered RescueMu transposon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GSS
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49 c 200 g 143 t
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                 /note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site_1: BamHI; Site_2: BglII; PBLUEScript backbone); BamHI; BamHI; Site_2: BglII; RescueMu is a 4.9 kb, modified maize Mu transposon designed to allow plasmid rescue from total genomic DNA.
                                                                                                       /clone_lib="1008 - RescueMu Grid
/tissue_type="leaf"
/dev_stage="adult"
                                                                                                                                                                /organism="Zea mays"
/cultivar="mixed background W23/A188/B73"
/db_xref="taxon:4577"
                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                         /lab_host="DH10B"
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I Zea mays genomic,
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 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Predni, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BH751161 112 bp DNA linear GSS SALK_049464.49.75.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK_049464.49.75.x,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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BH751161
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                                                                                                                                                                                                                                                                                                                                                                                                                     ecker@salk.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              units. For more information on RescueMu, go to the web site www.zmdb.iastate.edu and follow the links for 'RescueMu.' Grid I was grown at Berkeley in 2001. DNA was extracted from leaf punches, double digested using Bami and BglII, and ligated to form circular plasmids. DH10F cells were transformed and then screened on LB plates wi
                                                                                          directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://slgnal.salk.edu/tdna_protocols.html" 26 c 25 g 30 t
                                                                                                                                                                  /clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
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                                                                                                                                                                                                                                             /strain="Columbia 0"
/db_xref="taxon:3702"
/clone="SALK_049464.49.75.x"
                                                                                                                                                                                                                                                                                        /organism="Arabidopsis thaliana"
/strain="Columbia 0"
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Kim, C.J., Parker, H.,
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                                                                                                                                                       GSS
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ,C., Jeske,A., Karnes,M., Kim,Č.J., Parker,H., Prednis,L
Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
Arabidopsis Genome
Unpublished (2001)
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      Alonso, J.M.,
                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rossidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 153)
                                                                                                          Arabidopsis thaliana
                                                                                                                                 thale cress.
                                                                                                                                                                           BH619283.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:3702"
/clone="SALK_047736.48.15.x"
/clone="SALK_047736.48.15.x"
/clone="Parabidopsis thaliana TDNA insertion lines"
/clone="Parabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"
a 36 c 30 g 35 t
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Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., Gadrinab
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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6.2;
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Prednis, L., Shinn,
                                                                                                                                                                                                                                     lines Arabidopsis
                                                                                                                                                                                                                                                             GSS 30-JAN-2002
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BASE COUNT
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AUTHORS
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KEYWORDS
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DEFINITION
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BH747013
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Matches
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source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 296
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BH747013 153 bp SALK_008070.43.05.x Arabidopsis the Arabidopsis thaliana genomic clone
                                                                                                                                                                                                                                                                        ,C., Jeske,A., Karnes,M., Kim,Č.J., Parker,H., Predni
, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10010 N. Torrey Pines Road,
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                      Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                                                                                                                              Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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Arabidopsis thaliana
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
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                                   Class: TDNA tagged
                                                                                                                   Tel: 858 453 4100 x1752 Fax: 858 558 6379
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                                                                          This is single pass
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ecker@salk.edu
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                                                                                                                                                                                                                                                                                                                                                       (bases 1 to 153)
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larity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36
                                                                                                 ecker@salk.edu
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/Clone_lib="Arabidopsis thaliana lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 34 c 36 g 47 t
                Location/Qualifiers
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/db_xref="taxon:3702"
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/etrain="Columbia 0"
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                                                                            sequence
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Pred. No.
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                                                                            recovered
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6.3;
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                                                                                                                                                                                                                                                                                                                                Chen, H.,
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                                                                            from the
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                                                                            left border of
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lines
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ACCESSION
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BH748291/c
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ORIGIN
                                                                                                              BASE COUNT
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                                             Query Match
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Best Local 9
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      Matches
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A Sequence-Indexed Library of Insertion Mutations in the Arabidopsis Genome Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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вн748291.1
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                        Similarity
  9.9%;
ilarity 100.0%;
Conservative
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 35 c 37 g 49 t
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                                                                                                                                                                                                                                                                                                                                /strain="Columbia
                                                                                                                                                                                                                                                                                                                                                     organism-"Arabidopsis thaliana"
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/strain="Columbia 0"
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                      Score 39; DB 17;
Pred. No. 6.3;
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Pred. No.
    Mismatches
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6.3;
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SALK_016522.
Arabidopsis
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вн746616.1
                                                                                                              BH746616 175 bp DNA linear GSS SALK_045268.54.50.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK_045268.54.50.x,
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The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
Arabidopsis thaliana
                                       GSS
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/note="por was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 43 c 39 g 38 t
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/strain="Columbia 0"
/db_xref="taxon:3702"
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Spermatophyta; Magnollophyta; Liliopsida; Poales; Poaceae; PACC
clade; Panicoideae; Andropogoneae; Zea.
1 (bases 1 to 177)
Walbot,V.
Class: transposon-tagged
               Very probable ligation site found so sequence was trimmed Post-ligation sequence submitted separately. Plate: 1008026 row: 10
                                                                                                        Tel: 650 723 2227 Fax: 650 725 8221
                                                                                                                                                                                                                   Contact: Walbot V
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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Unpublished (2001)
                                                                                 Email: walbot@stanford.edu
                                                                                                                                                 855 California Ave, Palo Alto,
                                                                                                                                                                        Stanford University
                                                                                                                                                                                          Department of Biological Sciences
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A Sequence-Indexed Library of Insertion Mutations in the
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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AlonsO,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn, J. Zimmerman,J. and Ecker,J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; Spermatophyta; Spermatophyt
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39; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thale cress.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ecker@salk.edu
/clone="SALK_019366.54.25.x"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
                                                                                                                                                       /strain="Columbia 0"
/db_xref="taxon:3702"
                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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/dev starr"
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/strain="Columbia 0"
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/db_xref="taxon:4577"
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elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"

ORIGIN

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Query Match 9.9%; Score 39; DB 17; Length 190;
Best Local Similarity 100.0%; Pred. No. 6.4;
Matches 39; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Search completed: April 14, 2003, 10:25:42 Job time: 916.489 secs

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ALIGNMENTS

JOURNAL	TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AX036754	RESULT 1
mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 20 05-OCT-2000;	Chimeric expression promoters originating from commelina yellow	Rance, I., Theisen, M. and Gruber, V.	1 (bases 1 to 462)	artificial sequences.	synthetic construct	synthetic construct.	•	AX036754.1 GI:11226263	AX036754	Sequence 20 from Patent WO0058485.	AX036754 462 bp DNA linear		
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                                                       /organism="synthetic construct"
/db_xref="taxon:32630"
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                                             /note-"promoter MPr1165"
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Pred. No. 3.2e-116;
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Best Local S
Matches 385
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Best Local Sim
Matches 376;
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AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
                                               TGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGAGGAGATGAA 5858
                                                         TGCCTGCAGCACCTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGAGGAGATGAA 69
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385; Conserv
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                                                                                                                                                                                                                                                                                                                       synthetic construct artificial sequences.
1 (bases 1 to 9285)
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Sequence 52 from Patent
AX093047
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                                                                                                                                                                                                                                                                                                                                                          synthetic construct
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                                                                                            Conservative
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                                                                                                                                                 /note="pMRT1336 results from the insertion into pMRT1322 a of the promoter MPT1165 isolated from plasmid pMRT1322 a described in PCT patent application PCT/IB00/00370" 2252 c 2506 g 2087 t
                                                                                                                                                                                                        /organism="synthetic construct"
/db_xref="taxon:32630"
_note="pMRT1336"
                                                                                                                                                                                                                                           Location/Qualifiers
1. .9285
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95.28;
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Pred. No. 1.6e-89;
0; Mismatches 19
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GATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTAGT
                                                            AAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAA 189
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1 (bases 1 to 15077)
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Sequence 57 from Patent
AX093052
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                                                                                                                                                                                                                                                                        /organism="synthetic construct"
/db_xref="taxon:32630"
_note="pMRT1342"
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AX036758
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                                                                                                                                                                                                                                                                                                                   Local
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         ACCAACACAACAACCTAGAGGATCC 462
                                                                                                                            GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC 369
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                                                                                                                                                                                                                                                                                                         al Similarity
379; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                      artificial sequences.

1 (bases 1 to 541)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating 1
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 24 05-007-2000;
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Sequence
AX036758
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Pred. No. 1.3e-86;
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 Rance, I.,
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AX036753
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1 (bases 1 to 393)

Rance,I., Theisen,M. and Gruber,V.

Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 19 05-0CT-2000;
                   artificial sequences.
1 (bases 1 to 604)
                                      synthetic construct. synthetic construct
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, Theisen, M. and expression promo
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/db_xref="taxon:32630"
/note="promoter MPr1162"
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160 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA
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Ax036755.1
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                                                                                                      Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating i
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 21 05-OCT-2000;
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1 (bases 1 to 392)
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/db_xref="taxon:32630"
/note="promoter MPr1167"
            /organism="synthetic c
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/note="promoter MPr1164"
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                                                                                                                                                artificial sequences.

1 (bases 1 to 317)

Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0.058485-A 3 05-0CT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                   AX036737 317 bp
Sequence 3 from Patent WO0058485
AX036737
                                                                                                                                                                                                                              synthetic construct
                                                                                                                                                                                                                                        synthetic construct.
                                                                                                                                                                                                                                                          AX036737.1 GI:11226246
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                                       Conservative
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                                                                                   /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1116"
1. 317
61 c 74 g 75 t
                                                                                                                                          Location/Qualifiers
                                                46.0%;
99.5%;
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Pred. No. 2.8e
0; Mismatches
                                      0;
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                                               Score 212.4; DB 6
Pred. No. 1.2e-47;
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                                       Mismatches
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1. No. 2.8e-55;
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VERSION
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AX036738
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ACCESSION
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BASE COUNT
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                            AX036738
Sequence
AX036738
synthetic construct
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                   AX036738.1
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AX036759
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                   GI:11226247
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138 GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
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                                                                                                                                                       GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           artificial sequences.

1 (bases 1 to 472)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus
Patent: WO 0050485-A 25 05-0CT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
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1. .472
1. 92 c 112 g 11
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/db_xref="taxon:32630"
348 bp
Patent W00058485
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Pred. No. 1.2e
0; Mismatches
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WO0058485
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1.2e-47;
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TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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                                                             Query Match
Best Local
                                                 Matches 154;
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CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 189
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1 (bases 1 to 348)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus patent: WO 0050485-A 4 05-OCT-2000;

MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);

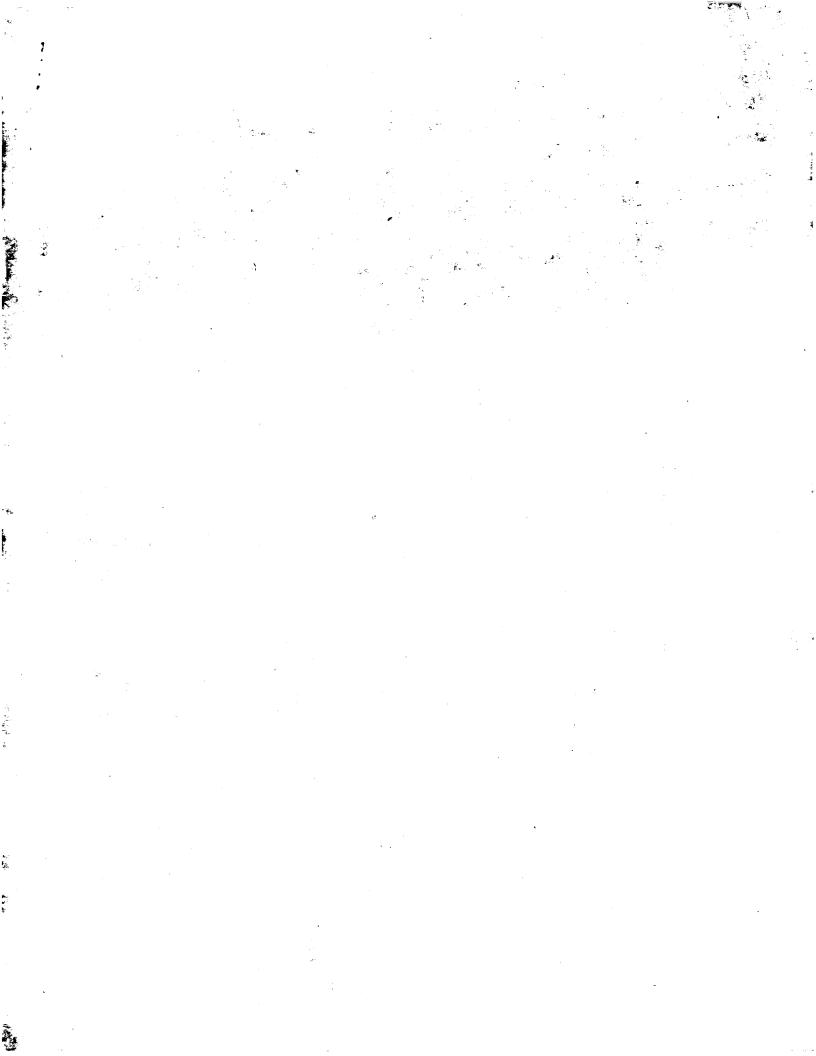
GRUBER VERONIQUE (FR);
                                                                                                                                                                                               Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 5 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                 synthetic construct. synthetic construct
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Sequence 5 from Patent WO0058485
AX036739
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l (bases 1 to 371)
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                                                           Similarity
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/db_xref="taxon:32630"
/note="promoter MPr1117"
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/db_xref="taxon:32630"
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                                                                                                                                    note="promoter MPr1146"
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                                                           Score 152.8;
Pred. No. 2.9
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REFERENCE
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BASE COUNT
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AX036741
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Best Local S
Matches 153
                                                                                                                       TITLE
                                                                                             JOURNAL
                                                                                                                                      AUTHORS
                                                                                                                                                                              ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                        GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 181
                                                                                                                                                                                                                                                                                                                                                                                                      GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197
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                                                                                                                                                                                                                                                                                                                                    GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
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                                                                                                                                                                                                                                   AX036740
Sequence 6 :
AX036740
                                                              Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 6 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 301)
Rance, I., Thetsen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus patent: WO 0058485-A 7 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); GRUBER VERONIQUE (FR)
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Ax036741
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1 (bases 1 to 398)
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/db_xref="taxon:32630"
                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="promoter MPr1154"
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/db_xref="taxon:32630"
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from Patent WO0058485
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Patent WO0058485.
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Pred. No. 5.7e-28;
0; Mismatches 2
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SOURCE
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AX036736
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Search completed: April 14, 2003, 09:29:25
Job time : 1330.64 secs
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                                                                                                                                                                                           Query Match 26.1%;
Best Local Similarity 93.3%;
Matches 126; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM
                                                                                                                                                                                                                                                                                                                                                                              JOURNAL
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                                              336 ATCACAAAGGAATCT 350
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                                                                                                                                                                                                                                                                                                                                              synthetic construct.
synthetic construct
artificial sequences.
1 (bases 1 to 515)
1 (bases 1 to 515)
Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 2 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR) GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AX036736 515 bp
Sequence 2 from Patent WO0058485.
AX036736
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                                                                                                                                                                                                                                                                         /organism="synthetic construct"
/db_xref="taxon:32630"
/note="Promoter from the intergenic region of Cassava Vein Mosair Virus of 515 bp in length EMBL U59751"
                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
1. .515
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80 c 93 g 97
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79 c
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Pred. No. 5.6e-28;
0; Mismatches 2;
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The present sequence represents a chimeric promoter of the invention. These specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived chimeric promoter comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression a second plant promoter comprising a plant promoter originates from a second plant promoter originates from the commelina yellow mottle virus, and the second plant promoter are commelina yellow mottle virus, and the second plant promoters are derived from intergenic regions. The chimeric promoters are useful derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 462 BP; 148 A; 87 C; 111 G; 116 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      121 AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACACATAGAGGACACATGTA 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     241 GATTGATGTGATATCAAGATTGATGTGATGTCTCCACTGACGTAAGGGATGACGCATGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANCANTGANANGANGANGATANGGTCGGTGATTGTGANAGAGACATAGAGGACACATGTA 180
                           Promoter; intergenic region; Commelina yellow mot Mpr1165. Chimeric expression promoter; plant yellow mot Mpr1165. plant green tissue expression promoter; cassava vein mipro transgenic plant; chimera; ss.
                                                                                                                                                                           Nucleotide sequence
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100.0%; Pred. No. 1e-131;
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                           in promoter;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chimeric expression promoter for transgenic plant production, comprise sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 600 BP; 188 A; 111 C; 147 G; 154 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for producing transgenic plants.
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Local 5
385;
                                                                                                                                                                                                                                                                                                                                                                                                                                            121 AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
|| 121 || 122 || 123 || 124 || 124 || 125 || 126 || 127 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 128 || 
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Cassava vein mosaic virus.
                                                                                                                                                                           Gruber V,
                                                                                                                                   ACGCATGCCACGTTACCCGGTATGCCGGTTCCCCAAGCTTTATTT
                                                                                          ACGCATGCCACGACTAGTGATTGATGTGATATCAAGATTGATGT
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95.3%;
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121 AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
121 AACAATGAAAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
121 AACAATGAAAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 121 AACAATGAAAAGAAGAAGATAAAGGTCGGTGATTGTGAAAGAGACATAGAGAGACATGTA
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181 AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGACTAGT 240
181 AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTT----- 231
181 AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTT----- 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                241 GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ncleotide sequence of chimeric expression promoter MP11167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ter; intergenic region; Commelina yellow mottle virus; to expression promoter; plant vascular expression mosaic virus; to expression promoter; Cassava vein mosaic virus; een tissue expression promoter; Cassava
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   397 ATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACCTAGTGAT 456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          433 AGAAAACCAACAACAACCTAGAAGGATCC 462
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h 83.3%;
                                                                                                                                                                                                                                                promoter; intergenic region; Commelina yellow mottle virus; promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; Cassava veln mosaic virus; plant green tissue expression promoter; transgenic plant; chimera; SS.
                                                                                                                                                                                                                                                                                                                                       Nucleotide sequence of chimeric expression promoter MPr1164.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric - Commelina yellow mottle virus.
Chimeric - Cassava veln mosaic virus.
                                                                                                                                                                WO200058485-A1.
                                                                                                                             05-OCT-2000
                                                                                         29-MAR-2000; 2000WO-IB00370.
                                                         29-MAR-1999;
                   (MERI-) MERISTEM THERAPEUTICS.
                                                                      99FR-0003925.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 262; pB 21; Length 604; pred. No. 2.7e-70;
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The present sequence represents a chimeric chimeric promoters comprise a nucleic expression a second plant promoter, in which a plant vascerity is replaced with a nucleic acid sequent served with a nucleic acid sequent sequention.

The present sequence represents a chimeric expression is replaced with a nucleic acid sequent invention. The promoter comprising a plant order vasceries in the sequential promoter comprising a plant order result is derived to the present the present of the present comprision and the present comprision

Chimeric expression provided in the sequence from promoter current sequence from promoter

Page 87; 91pp; English.

(Mava vein mosaic virus. mmelina yellow mottle virus.

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RESULT 7
AAA96837
ID AAA9
XX
AC AAA9
XX
DT 19-F
XX
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XX
KW Prom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
Promoter; intergenic region; Commelina yellow mottle virus;
                          Nucleotide sequence of chimeric expression promoter MPr1116
                                                                                                                  AAA96837
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               derived from intergenic regions. The chimeric promoters are useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 AAGCTTGCATGCCTGCAGCACCTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA 60
                                                                                                                                                                                                  GTGTAGTAGCTTAGAAAACCAACAACAAC 451
                                                                                                                                                                                                                                                                                                                                                                                                                ACGCATGCCACGTTACCCGGTATGCCGGTTCCCCAAGCTTTATTTCCCTTATTTAAGCACTT 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC 120
                                                                                                                                                                                    GTGTAGTAGCTTAGAAAACCAACACAACAAC 382
                                                                                                                                                                                                                                           AATCACTAGTCGTTACCCGGTATGCCGGTTCCCCAAGCTTTATTTCCTTATTTAAGCACTT
                                                                                                                                                                                                                                                                                                 TTACGTCAGTGGAGATATCACATCAATCTTGATATCACATC---
                                                                                                                                                                                                                                                                                                                 ACGACTAGTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATG
                                                                                                                                                                                                                                                                                                                                                                                    GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGCTTGCATGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                349;
                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page
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                                                        (first entry)
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                                                                                                                DNA;
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77.4%;
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Pred. No. 1.1e-63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 21;
                                                                                                                                                                                                                                                                                                                                                         ----TGGCATGCGTCATCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              33; Indels - 69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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RESULT 8

AAA96859

standard; DNA; 472

B 5

200 198

233

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XEXTXAXI

19-FEB-2001

(first entry)

Nucleotide sequence of chimeric

expression

promoter

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Дb
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                                                                                                                                    Qy
                                                                                                                                                                                                                             Matches
                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                               The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commeilna yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric - Commelina yellow mottle vichimeric - Cassava vein mosaic virus
                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERI-)
                                                                                                                                                                                                                                                      for producing transgenic plants.
 138
                                                       78
                                                                                  80
                                                                                                              18
                                                                                                                                        20
                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 eric expression promoter for transgenic plant production, comprisence from promoter comprising vascular expression region replaced sequence from promoter comprising green tissue expression region
 TAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC
                TAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC
                                                      TGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGA
                                                                      2000-647238/62
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                                                                                                                                                                    al Similarity
213; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MERISTEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                           Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Commelina yellow mottle virus
                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                            81;
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                                                                                                                                                                                                                            107 A; 61 C;
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                                                                                                                                                                                   46.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Theisen M;
                                                                                                                                                                    0;
                                                                                                                                                                    Score 212.4; DB 21; pred. No. 3.4e-55; 0; Mismatches 1;
                                                                                                                                                                                                                            74 G; 75 T;
                                                                                                                                                                                                                            0 other;
                                                                                                                                                                    Indels
                                                                                                                                                                                             Length
                                                                                                                                                                    0
                                                                                                                                                                    Gaps
 197
                           199
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                                                                                                              77
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RESULT 9
AAA96838
ID AAA9
XX
AC AAA9
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                                                                                                                                                                                                                                                                                                                             В
                                                                                                                                                                                                                                                                                                                                                                                                                                              δ
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from commending yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric expression promoter for transgenic plant production, esequence from promoter comprising vascular expression region rewith sequence from promoter comprising green tissue expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric - Commelina yellow mottle virus. Chimeric - Cassava vein mosaic virus.
                             AAA96838 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Promotter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 5; Page 88; 91pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-OCT-2000
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                                                                                                   399
                                                                                                                                                                                                                  279
                                                                                                                                                                                                                                                                          220
                                                                                                                                                                                                                                                                                                    138
                                                                                                                                                                                                                                                                                                                                160
                                                                                                                                                                                                                                                                                                                                                          78
                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity es 236; Conserv
                                                                                                   GGTTCCCAAGCTTTATTT
                                                                                                                              TGATATCAAGATTGATGT 335
                                                                                                                                                        GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACCCGGTATGCT
                                                                                                                                                                                                             GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                              CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA
                                                                                                                                                                                                                                                                                                                                          CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000-647238/62
                                                                                                                                                                                   GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                              472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 149 A; 92 C; 112 G; 119 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99FR-0003925.
                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                  46.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Theisen M;
                                                                                                  416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                              348
                              ΒP
                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                 Score 212.4; DB Pred. No. 4e-55;
                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                  DВ
                                                                                                                                                                                                                                                                                                                                                                                     21;
                                                                                                                                                                                                                                                                                                                                                                                                                 21;
                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                               Length 472;
                                                                                                                                                                                                                                                                                                                                                                                     1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              replaced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  region
                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                 197
                                                                                                                                                                                                                                                                                                                             219
                                                                                                                                                                                                                                            257
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RESULT 10
AAA96839
ID AAA96
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Best Local S
                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric
                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transgenic plant; chimera; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAR-1999;
                                                                                                                                 172
                                                                                                                                                                                   112
                                                                              231
                                                                                                        200
                                                                                                                                                        140
                                                                                                                                                                                                                                                                                        Local Sir
hes 212;
                                                                                                                                                                                                           80
                                                                                                                                                                                                                                      52
                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present sequence represents a chimeric promoter of the invention.
                                                                                                                                                                                                                                                                                                                                                                    producing
                                                                                                                             GGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
                                                                                                     GGAAAGTAACCTTATGCATTTGTAATTTGGTTAC
                                                                                                                                                                                   TGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGA
                                                                                                                                                                                                           TGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGA 139
                                                                                                                                                                                                                                     ric expression promoter; green tissue expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5; Page 81; 91pp; English
                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                            348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               intergenic region; Commelina yellow mottle virus; expression promoter; plant vascular expression proen tissue expression promoter; Cassava vein mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Commelina yellow
Cassava vein mosa
                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                          BP; 116
                                                                                                                                                                                                                                                                                                                                                                    transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99FR-0003925
                                                                                                                                                                                                                                                                                                     43.48;
99.18;
                                                                                                                                                                                                                                                                                                                                          A; 70 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mosaic
                                                                                                                                                                                                                                                                                                                                                                    plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mottle virus
aic virus.
                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                   Score 200.4; DB 2
Pred. No. 1.7e-51;
                                                                                                                                                                                                                                                                                                                                            78
                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                            <u>.</u>
                                                                                                                                                                                                                                                                                                                                         84 T; 0 other;
                                                                            264
                                                                                                                                                                                                                                                                                                                 DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   promoter
                                                                                                                                                                                                                                                                                                                 Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              promoter;
                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                      111
                                                                                                                                                                                   171
                                                                                                                                                         199
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AAA96839

standard;

DNA;

371

ВР

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RESULT 11
AAA96841
ID AAA96
XX
AC AAA96
XX
DT 19-FE
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                                                                                                                                                                               В
                                                                                                                                                                                                Qy
                                                                                                                                                                                                                                                                 Matches 154;
                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                 The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric - Commelia
Chimeric - Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence from with sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
 19-FEB-2001
                      AAA96841;
                                          AAA96841 standard; DNA;
                                                                                                                                                                                                                                                                 Sequence 371 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-647238/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleotide sequence of chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-FEB-2001
                                                                                                                                       190
                                                                                                                                                           138
                                                                                                                                                                                130
                                                                                                                 198
                                                                                                                                                                                         78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACGACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                   Local
                                                                                            GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
                                                                                                                                               GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197
                                                                                                                                     GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                              CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 189
                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            expression promoter for transgenic plant production, comprise from promoter comprising vascular expression region replaced uence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                 Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Commelina yellow mottle virus.
                                                                                                                                                                                                                         Conservative
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                 81; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99FR-0003925
                                                                                                                                                                                                                                                                122 A; 68 C; 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vein mosaic virus
                                                                                                                                                                                                                                33.1%;
98.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Theisen M;
                                          301 BP
                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                       Score 152.8;
Pred. No. 7.1e
0; Mismatches
                                                                                                                                                                                                                                                                <u>.</u>
                                                                                                                                                                                                                                                                92 T; 0 other;
                                                                                                                                                                                                                                3; DB 21;
1.1e-37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            promoter MPr1146
                                                                                                                                                                                                                        2;
                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprises
                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                       0
                                                                                RESULT 12
AAA96840
XEXEXEX
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Nucleotide sequence of chimeric expression

promoter

19-FEB-2001

(first entry)

AAA96840 standard; DNA; 398

182

GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233

217

123 138

63 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA

122

;;

GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197

GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG

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Matches
                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric
Chimeric
                                                                                                                                                                                                                        The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rance I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleotide sequence of chimeric expression
                                                                                                                                                                                                            for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MERI-)
78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               eric expression promoter for transgenic plant production, comprisence from promoter comprising vascular expression region replaced sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-647238/62.
                                                                       153;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>ن</u>
                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MERISTEM THERAPEUTICS
                                                                                                                                                                  301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Commelina yellow mottle virus Cassava vein mosaic virus.
                                                                    Conservative
                                                                                                                                                               BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99FR-0003925
                                                                                                                                                             98 A; 54 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               91pp; English.
                                                                                        30.5%;
98.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Theisen
                                                                    0;
                                                               Score 140.8; DB 21; Pred. No. 3.1e-33; 0; Mismatches 2;
                                                                                                                                                               74 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Z
                                                                                                                                                               75 T; 0
                                                                                                              DB 21;
                                                                                                                                                               other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         promoter MPr1154
                                                                    Indels
                                                                                                                 Length
                                                                                                                 301;
                                                                  1;
                                                                  Gaps
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ARESULT 13
AAV14019
ID AAV14
XX AV14
AC AAV14
XX 18-JU
DT 18-JU
XX CSVMV
XX CSVMV
XX CASSA
KW tissu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ð
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity Matches 153; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                             Cassava vein mosaic virus; CsVMV; promoter; cultivated crop; tissue-specific expression control; transgenic plant; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
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Chimeric - Cassava
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                                                                                                                        CsVMV promoter CVP1
                                                                                                                                                                                         18-JUN-1998
                                                                                                                                                                                                                                                                                                                     AAV14019 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 398 BP; 128 A; 80 C; 93 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 5; Page 82; 91pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GATAAGGTCGGTGATTGTGAAAGAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Commelina yellow mottle virus. Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 140.8;
Pred. No. 3.5e
0; Mismatches
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2;
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RESULT 14
AAV14021
ID AAV14
XX AAV14
XX AAV14
XX AAV14
XX CASSA
KW CASSA
KW CASSA
KW CASSA
KW CASSA
KW CASSA
XX CASSA
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XX WO974
XX YO974
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Best Local Similarity
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                                                                                                                      Cassava vein mosaic virus; CsVMV; promoter; cultivated tissue-specific expression control; transgenic plant; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cassava vein mosaic virus promoter - used to express heterologous DNA sequences for producing transgenic plants having altered
                                                                                                                                                                  CsVMV promoter pB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Page 74; 115pp;
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                                                                                                                                                                                                                               AAV14021;
                                                                                                                                                                                                                                                           AAV14021 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                 153 TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93 GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT 152
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                                                                                                                                                                                                                                                                                                                                                                   ATGCATTTGTAATTT
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                                                                                           mosaic
                                                                                                                                                                                               (first entry)
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93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                           ВP
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Pred. No. 5.5e-27;
0; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                83 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Verdaguer
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24-DEC-1997

20-JUN-1997;

97WO-US10376

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RESULT 15
AAA96836
XX AAA96
XX AAA96
XX 19-FE
XX Promo
XX Promo
XW Chime
KW Chime
KW Chime
KW Clant
XW Cassa
XX WO20C
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Best Local S
Matches 126
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                                                                                                                                                                                                                                                                                                                                                                             Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA96836 standard; DNA;
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                            (MERI-) MERISTEM THERAPEUTICS
                                                                               29-MAR-1999;
                                                                                                                                  29-MAR-2000;
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                                                                                                                                                                                                                                                                                                   Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                         transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Promoter from intergenic region of Cassava vein mosaic virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT 212
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                                                                                                                                                                                                                                                                                                                                                         plant; ss.
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                                                                                                                                     2000WO-IB00370
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0; Mismatches
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                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 126; Conserv
                                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents a promoter fragment from the intergenic region of Cassava vein mosaic virus. The promoter is used to construct chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                      Sequence 515 BP; 198 A; 79 C; 109 G; 129 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                    The chimeric promoters are useful for producing transgenic plants
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93.3%;
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Sequence Sequence Patent No.

30, Appl 30, Appl . 5254799

Sequence Sequence

Sequence Seq

46, Appl 43, Appl 85, Appl 8, Appli 46, Appli 46, Appli 2, Appli 2, Appli 5, Appli 5, Appli 5, Appli 5, Appli 17, Appli

Sequence

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/FCTUS_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
       GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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 US-08-232-463-14
US-08-247-809A-3
US-08-711-72B-3
US-08-764-100-23
US-09-764-100-23
US-09-330-760-1
US-09-330-737-1
US-09-330-714A-1
US-09-330-714A-1
US-09-328-826-1
US-09-328-826-1
US-09-328-473-5
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US-09-330-714A-5
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US-09-328-473-5
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US-09-328-826-5
US-09-330-714A-5
US-08-31-746-1
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 Query Match
                                                                                                                                                                        APPLICATION NUMBER: US/07/935,313
FILING DATE:
APPLICATION NUMBER: EP 91 114 300.6
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REGISTRATION NUMBER: 30472/114 IMPLEDED ON THE COMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEPAX: (703)883-4109
                                                                                                                  TELEX: 899149
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 22313-0299

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: VA
COUNTRY: USA
COUNTRY: 22313-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
                                           IMMEDIATE SOURCE:
CLONE: pTZgpt-
                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                        TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: lin
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                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/232,463
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                                           pTZgpt-Fls
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1800 Diagonal Road, Suite 500
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US-08-729-601A-43
US-09-35-3-970-35
US-09-011-151-8
US-09-011-151-9
US-09-018-244-2
US-09-471-913-6
US-09-478-993-2
US-09-458-093-2
US-09-65-999-6
US-09-65-999-6
US-08-247-809A-5
US-08-247-809A-5
US-08-711-728-5
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US-08-463-308-30
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   Score 48.6;
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Length 7218;
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Result

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Score

Database :

Minimum DB Maximum DB

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Title:

Run on: OM nucleic -

Scoring table: Sequence: Perfect score:

Best Local Similarity Matches 18; Conserv

Conservative

Pred. No. 2.7e-05; 85; Mismatches 134;

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; MOLECULE TYPE: US-08-247-809A-3
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                                                                                                   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1247
                                                                                                                                                                    REGISTRATION NUMBER: 33,141
REFERENCE/DOCKET NUMBER: BAY
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 332-1700
                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 2.0 MB
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: DEOXYRIBONUCLEIC ACIDS NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                     APPLICATION NUMBER: P 43
FILING DATE: May 28, 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 800 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1067 RATCGCAAGCTCCCTCGACCTGCAGCCAAGCTCGGAA 1031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Peter H. Schreier; Klaus Stenzel; Gunter Adam;
APPLICANT: Edgar Maiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  359 TGACGCATGCCACGTTACCCGGTATGCCGGTTCCCAA 395
                                   TOPOLOGY:
                                               STRANDEDNESS:
                                                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
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CITY: Tarrytown
STATE: New York
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                                                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08 FILING DATE: May 23, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTGATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'RY: U.S.A.
10591-5144
                                                               nucleic acid
                                                                                                                                                                                                                                       Kurt G. Briscoe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: SPRUNG HORN KRAMER & WOODS 660 White Plains Road
                                                                                                                                                      (914) 332-1844
                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                           NEC Powermate 1 Plus
               DNA (genomic)
                                                  single
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RESULT 4
US-08-764-100-23
Sequence 23, Application US/08764100
; Patent No. 5773700
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US-08-711-728-3
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Best Local Similarity 88.u
                                                                                                                                                                      Query Match
Best Local :
                                                                                                                                                        Matches
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 23-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION UNMBER: DE 4:
FILING DATE: 28-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (914) 332-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                              338 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  338 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 387
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                                                                                                                           258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/711,728 FILING DATE: 03-SEPT-1996
                                                                                                                                                                    Local
                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                            TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Kurt G. Briscoe
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3, Application US/08711728
5, 5973135
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                                                                                                                                                                    Similarity
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Edgar Maiss
                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                             linear
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Pred. No. 0.0028;
0; Mismatches 6;
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Pred. No. 0.0028;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/214,064
FILING DATE:
APPLICATION NUMBER: US 08/032,235
FILING DATE: 17-MAR-1993
APPLICATION NUMBER: GB 9206016.9
FILING DATE: 19-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO. 5773700r1s, Allen E.
REGISTRATION NUMBER: 34,490
REFERENCE/DOCKET NUMBER: 137-1061
                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 857-1125 INFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                             NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6114608artis Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/764,100
FILLING DATE: 06-DEC-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 446 base pairs
                                                                                                                                                                                                    APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                      274 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 323
                                                                                           CITY: Summit
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
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ZIP: 94304
                                                                                                                               STREET:
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STREET: 975 California Avenue
                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                           07901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CA
                                                                                                                             564 Morris Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                            USA
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Pred. No. 0.0028;
0; Mismatches 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6222104
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                                   TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/0
FILING DATE: March 13, 1
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
TYPE: nucleic acid
                  SEQUENCE CHARACTERISTICS:
                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 62221
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ANTI-SENSE: N
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                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                        REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 13
                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Summit
STATE: New Jersey
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Local Similarity 88.0%;
les 44; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/042,426 FILING DATE: March 13, 1998
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LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                07901
                                                                                                                                           Hoxie, Thomas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   564 Morris Avenue
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                                                        Matches
                                                                                   Query Match
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Patent No. 6229075
GENERAL INFORMATION:
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                                                                                                                          ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: 35S Promoter
                                                                                                                                                       HYPOTHETICAL: P
                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 13:
REFERENCE/DOCKET NUMBER: 13:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/042,426
FILING DATE: March 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
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CORRESPONDENCE ADDRESS:
ADDRESSEE: NO. 6229075artis Corporation
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CLONE: 35S Promoter
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                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: DNA
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TITLE OF IN
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 274
             258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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CITY: Summit
                                                                      Local
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                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 323
                                                                      Similarity
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                                                      8.7%;
llarity 88.0%;
Conservative
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ENTION: DNA Construct Containing Bacillus
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                                                      Score 40.4; DB Pred. No. 0.003; Mismatches
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                                                                                DB 4;
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                                                                              Length 532;
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RESULT 9
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                                                                                                                                                                                              Sequence 1, Application US/09330737 Patent No. 6232534
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                                                                                           NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6232534art1s Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                            ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: 35S Promoter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPAX: (919) 541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/042,426
FILING DATE: March 13, 1998
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy
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CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6232533art1s Corporation
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                                                                                                                                                TITLE OF INVENTION: DNA Construct
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                 274 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 323
                                                                                                                                                                                                                                                                                                                  258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                                              CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
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STATE: New Jersey
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                            COUNTRY:
                                                                               STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Hoxie, Thomas REGISTRATION NUMBER:
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                                             New Jersey
                                                                            564 Morris Avenue
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                              USA
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                                                                                                                                                               Irvin J.
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DNA Construct Containing Bacillus
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                                                                                                                                                               Mettler, Paul
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                                                                                                                                             S. Dietrich, Ralph Sinibaldi
Containing Bacillus
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OPERATING SYSTEM: PC-DOS/MS-DOS

IBM PC compatible

Floppy disk

MEDIUM TYPE:

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RESULT 10
US-09-329-169-1
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Matches 44; Conserv
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Patent No. 6329575
GENERAL INFORMATION:
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CLONE: 35S Promoter
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MOLECULE TYPE: DN
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REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 131
TELECOMMUNICATION INFORMATION:
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ETLING DATE: March 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Hoxie, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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TOPOLOGY: linear
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                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-TOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                      APPLICATION NUMBER: US/09/329,169
FILING DATE: 09-Jun-1999
CLASSIFICATION: - Unknown>
PRIOR APPLICATION DATA:
                                                                           APPLICATION NUMBER: 09/042,426 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6329575artis Corporation
STREET: 564 Morris Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
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                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 07901
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                                                                                                                                                                                                                                                                                                                                                                      STATE: New Jersey
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ilarity 88.0%;
Conservative
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US-09-330-714A-1
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Best Local (
                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
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                                                                                                                                                                                                                                          NAME: HOXIE, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 13:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/330,714A
FILING DATE: 11-Jun-1999
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/042,426
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
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TITLE OF INVENTION: DNA Construct Containing Bacillus
Thuringiensis Gene
              CLONE: 35S Promoter SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                      IMMEDIATE SOURCE:
CLONE: 35S P
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STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                        ANTI-SENSE:
                                                                                            HYPOTHETICAL:
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CLONE: 35S P
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ANTI-SENSE: NO
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                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: double
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STATE: New Jersey
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RESULT 13
US-09-042-426-5
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US-09-328-826-1
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                                                  Sequence 5, Application US/09042426 Patent No. 6114608
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Best Local
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Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/042,426
FILING DATE: MARCH 13, 1998
ATTORNEY/ACENT INFORMATION:
NAME: HOX16, Thomas
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                             CLONE: 35S Promoter SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
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TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
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ADDRESSEE: No. 6399860artis Corporation
                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 532 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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FILING DATE: 09-Jun-1999
CLASSIFICATION: <Unknown>
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 13
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                                                                                                                                                                                                                             Score 40.4; DB 4; Pred. No. 0.003;
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RESULT 14
US-09-291-238-5
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                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5, Application US/09291238 Patent No. 6222104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 5:
                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE
HYPOTHETICAL: N
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09,
FILING DATE: March 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
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ADDRESSEE: No. 6114608artis Corporation
                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                 STREET: ...
STREET: ...
STREET: ...
STREET: ...
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               322 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                                                                                                                              COUNTRY: UZIP: 07901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/0 FILING DATE: March 13, 1998 CLASSIFICATION: 800
                                                                     FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Hoxie, Thomas REGISTRATION NUMBER: 32,993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: UZIP: 07901
                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                       Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
VENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                    No. 6222104artis Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Floppy disk
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   09/042,426
13, 1998
                                                                                     US/09/291,238
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ATTORNEY/AGENT INFORMATION:

Hoxie, Thomas

135/1

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RESULT 15
US-09-330-760-5
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; CLONE: 35S Promoter
US-09-291-238-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5, Application US/09330760 Patent No. 6229075
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Best Local Similarity 88.0%;
Matches 44; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Irvin
                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/042,426
FILING DATE: MAICH 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOXIE, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (319) 541-8614
TELEPHONE: (319) 541-8614
                                                                                                                                            TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                LENGTH: 560 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomi
                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/330,760
FILING DATE:
CTARCOTTARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA CONSTRUCT CONTAINING Bacillus
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                  HYPOTHETICAL: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MMEDIATE SOURCE:
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CITY: Summit
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TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               564 Morris Avenue
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                                                       DNA (genomic)
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; CLONE: US-09-330-760-5
             35s Promoter
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Query Match 8.7 Best Local Similarity 88.0 Matches 44; Conservative 8.7%; Score 40.4; DB 4; Length 560; 88.0%; Pred. No. 0.0031; ative 0; Mismatches 6; Indels 0,

Gaps

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258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307

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Search completed: April 14, 2003, 10:28:36
Job time: 53.6289 secs

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Published_Applications_NA: *
/cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US07_REW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_REW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US08_REW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_REW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_REW_PUB.seq:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (without alignments)
7293.024 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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18	15 16	12 13	10 11	4005	321	0.
75.4	120.6 120.6	140.8 120.6 120.6	200.4 152.8 140.8	314 262 240.2 212.4 212.4	462 373.6 354.2	Score
17.3	26.1 26.1	30.5 26.1 26.1	33.1	58.0 52.0 46.0	100.0 80.9 76.7	Query Match Length
243 392	8340 8340	398 515 532	348 371 301	393 604 392 317	462 600 541	Length
991	1010	999	9999	وووو	إووو	DB
US-09-963-803-21 US-09-963-803-21	US-09-847-057-4 US-09-874-926-4 US-10-033-190-5	US-09-963-803-6 US-09-963-803-2 US-09-765-555-1	US-09-963-803-2 US-09-963-803-5 US-09-963-803-7	US-09-963-803-19 US-09-963-803-23 US-09-963-803-21 US-09-963-803-3	US-09-963-803-20 US-09-963-803-22 US-09-963-803-24	ID
Sequence 1, Appli Sequence 21, Appli Sequence 21, Appl	л 4- 4-	1,26	Sequence 2, Appli Sequence 4, Appli Sequence 5, Appli Sequence 7, Appli	19, 23, 21,	Sequence 20, Appl Sequence 22, Appl Sequence 24, Appl	Description

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5	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	
39.2	39.2	39.2	40.4	40.4	40.4	40.4	40.4	40.4	40.4	40.4	40.4	40.4	41.6	45	55	55	58	58	58	58	58	60	60	62	
8.5	8.5	8.5	8.7	8.7	8.7	8.7	8.7	8.7	8.7	8.7	8.7	8.7	9.0	9.7	11.9	11.9	12.6	12.6	12.6	12.6	12.6	13.0	13.0	13.4	
282	219	63	14446	13737	7129	5767	4973	3983	763	700	661	566	671	63	08	65	472	332	299	236	80	63	60	62	
10	10	10	9	9	9	9	9	10	9	9	10	10	9	9	10				10	10	10	9	9	9	
us-09-870-375-9	US-09-870-375-8	US-09-870-375-33	US-09-810-861B-4	US-10-074-279-10	US-10-047-542-101	US-09-810-861B-3	US-09-990-659A-15	US-09-758-987-1	US-10-162-214-9	US-10-138-221-9	US-09-943-692-33	US-09-951-470-3	US-10-184-644-346	US-09-963-803-13	US-09-870-375-28	US-09-963-803-8	US-09-870-375-5	US-09-870-375-7	US-09-870-375-36	US-09-870-375-35	US-09-870-375-34	US-09-963-803-12	US-09-963-803-9	US-09-963-803-10	
Sequence 9, Appli	Sequence 8, Appli	Sequence 33, Appl	Sequence 4, Appli	Sequence 10, Appl	Sequence 101, App	Sequence 3, Appli	Sequence 15, Appl	Sequence 1, Appli	Sequence 9, Appli	Sequence 9, Appli	Sequence 33, Appl	Sequence 3, Appli	Sequence 346, App	Sequence 13, Appl	Sequence 28, Appl	Sequence 8, Appli	Sequence 5, Appli	Sequence 7, Appli	Sequence 36, Appl	Sequence 35, Appl	Sequence 34, Appl	Sequence 12, Appl	Sequence 9, Appli	Sequence 10, Appl	- 11

ALIGNMENTS

RESULT 1 US-09-963-803-20

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; LOCATION: (1)..(462)
; OTHER INFORMATION:
US-09-963-803-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 20
LENGTH: 462
                                                                                                                                                                                                                          Query Match 100.0%; Score 462; DB 9; Best Local Similarity 100.0%; Pred. No: 5.8e-129; Matches 462; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20, Application US/09963803 Publication No. US20030028922A1
                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: promoter MPr1163 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence FEATURE:
121 AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
                                         Length 462;
                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                       Gaps
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Sequence 22, Application US/09963803

Publication No. US20030028922A1

GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
ITILE OF INVENTION: Chimeric expression promoters originating from the following formula and caseava voin mosaic virus
FILE REFERENCE: 184332042

CURRENT FILING DATE: US/09/963,803

CURRENT FILING DATE: 1909-03-26

PRIOR APPLICATION NUMBER: PGT 1800/00370

PRIOR FILING DATE: 2000-10-05

PRIOR FILING DATE: 2000-10-05

PRIOR FILING DATE: 2000-10-05

PRIOR FILING DATE: 399

SOFTWARE: Patentin version 3.1

SEQ ID NO 22

LENGTH: 600

TYPE: DNA

ORGANISM: Artificial Sequence
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US-09-963-803-22
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OTHER INFORMATION: pro
FEATURE:
NAME/KEY: promoter
LOCATION: (1)..(600)
OTHER INFORMATION:
IS-09-963-803-22
                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 385; Conserv
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           AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGACTAGT
                                                                                  AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
                                                                                                                                       GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC
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                                                                   AACAATGAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
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                                                                                                                                                                                                                                     Score 373.6; DB Pred. No. 2.2e-100; Mismatches 1
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2.2e-102;
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; NAME/KEY: promoter
; LOCATION: (1)..(541)
; OTHER INFORMATION:
US-09-963-803-24
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US-09-963-803-24
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APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILLING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PCT 1800/00370
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEO ID NOS: 39
COUNTRIANCE: Date: Date: Date: 000-10-05
NUMBER OF SEO ID NOS: 39
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Matches
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SEQ ID NO 24
LENGTH: 541
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                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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OTHER INFORMATION: promoter MPr1168
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517
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                                                                CGGTATGCCGGTTCCCAAGCTTTATTTCCTTATTTAAGCACTTGTGTAGTAGCTTAGAAA 437
                                                                                                                       TGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACC
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                           ACCAACACAACCTAGAGGATCC
                                                     CGGTATGCTGGTTCCCAAGCTTTATTTCCTTATTTAA--ACTTGTGTAGTAGCTTAGAAA
                                                                                                            TGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACC
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379; Conserv
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98.48;
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Pred. No. 1.4e-96
O; Mismatches
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RESULT 5
US-09-63-803-23
; Sequence 23, Application US/09963803
; Publication No. US20030028922A1
; GENERAL INFORMATION:
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; LOCATION: (1)..(393)
; OTHER INFORMATION:
US-09-963-803-19
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US-09-963-803-19
Sequence 19, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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LENGTH: 393
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Best Local Similarity
Matches 393; Conserv
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
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PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT 1B00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Chimeric expression promoters originating TITLE OF INVENTION: virus and cassava vein mosaic virus
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                                                                                                                      GTGTAGTAGCTTAGAAAACCAACACCAACAACCTAGAGGATCC 393
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NAME/KEY: promoter : LOCATION: (1)..(604) : OTHER INFORMATION: US-09-963-803-23
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Sequence 21, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
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Best Local (
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TITLE OF INVENTION: Chimeric expression promoters origine
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PRIOR A
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NUMBER OF SEQ ID NOS: 39
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: promoter MPrl167
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                  575
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les 375; Conserv
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83.3%;
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Pred. No. 7.1e-69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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PRIOR APPLICATION NUMBER: FR 99/03925

from

commelina

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; PRIOR FILING DATE: 1999-03-29; PRIOR APPLICATION NUMBER: PCT IB00/00370; PRIOR FILING DATE: 2000-10-05; NUMBER OF SEQ ID NOS: 39; SOFTWARE: PatentIn version 3.1; SEQ ID NO 21; LENGTH: 392; TYPE: DNA; Artificial Sequence; PRIORIES.
                                                                                                                                                                                                                                                                                                                                                                                        В
                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q
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                                                                                                                                                                                                                                  Sequence 3, Application US/09963803 Publication No. US20030028922A1
                                                                                                                                                                                                                  GENERAL INFORMATION:
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Matches 349;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
               SEQ ID
                                                                       APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
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NAME/KEY: promoter
LOCATION: (1)..(392)
OTHER INFORMATION:
                                           PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                              SOFTWARE: PatentIn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: promoter MPr1164
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                                                                                                                                                                                                                                                                                                                                                          GTGTAGTAGCTTAGAAAACCAACAACAACAAC 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTACGTCAGTGGAGATATCACATCAATCTTGATATCACATC-----
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317
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                              version 3.1
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77.4%;
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Pred. No. 2e
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FEATURE:

NAME/KEY: promoter

LOCATION: (1)..(472)

OTHER INFORMATION:
US-09-963-803-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 25, Application US/09963803 Publication No. US20030028922A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1
SEQ ID NO 25
LENGTH: 472
                                                                                                                                                                        Best
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/99/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 2000-10-05
RIOR FILING DATE: 2000-10-05
RUMBER: OF SEQ ID NOS: 39
                                                                                                                                                        Matches
                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                            OTHER INFORMATION: promoter MPr1169
                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION:
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GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
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                                                                                                                                                        Conservative
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                                                                                                                                                      Score 212.4; DB 9;
Pred. No. 4.8e-54;
0; Mismatches 21;
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Pred. No. 4e-
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                                                                                                                                                                                     Length 472;
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; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(348)
; OTHER INFORMATION:
US-09-963-803-4
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Sequence 5, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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SEQ ID NO 4
LENGTH: 348
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CURRENT FILLING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
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OTHER INFORMATION: promoter MPr1117
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                                                                                                                                                                                                                                                                                                                                                                           200 GGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 43.4%;
Local Similarity 99.1%;
es 212; Conservative
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Qy Db

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138 GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197

CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 122

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Publication No. US20030028922A1
GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 5
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Best Local
                                                    Matches
                                                                  Query Match
Best Local
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
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                                                                                                                                                                          FEATURE: promoter MPr1154
FEATURE:
                                                                                                                                     NAME/KEY: promoter LOCATION: (1)..(301) OTHER INFORMATION:
                                                                                                                                                                                                                                           LENGTH: 301
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
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                                                                  Similarity
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                                                  Conservative
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                                              Score 140.8; DB 9 
Pred. No. 1.2e-32; 
0; Mismatches 2
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Pred. No. 3.4
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Sequence 2, Application US/09963803

Publication No. US20030028922A1

GENERAL INCORNATION:

APPLICANT: Meristem Therapuetics

ITITLE OF INVENTION: Chineric expression promoters originating from the properties of INVENTION: Virus and cassava vein mosaic virus

FILE REFERENCE: 184332042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT FILING DATE: 2001-09-26

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR APPLICATION NUMBER: PRIOR OF SEQ ID NOS: 39

COUNTAIN DATE: 2000-10-05

PRIOR FILING DATE: 2000-10-05

PRIOR FILING DATE: 2000-10-05

PRIOR FILING DATE: 2000-10-05

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US-09-963-803-6
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                     SOFTWARE: PatentIn version 3.1 SEQ ID NO 2
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
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OTHER INFORMATION: promoter MPr1147
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es 153; Conserv
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RESULT 15
US-09-847-057-4/c
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Best Local
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; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(515)
; OTHER INFORMATION:
US-09-963-803-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: The Scripps Research Institute
TITLE OF INVENTION: Methods and compositions to
TITLE OF INVENTION: expression in plants
FILE REFERENCE: 27801-20014.40
CURRENT APPLICATION NUMBER: US/09/765,555
CURRENT APPLICATION NUMBER: US 09/620,897
PRIOR APPLICATION NUMBER: US 09/620,897
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: US 60/177,468
PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 75
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Promoter CSVMV US-09-765-555-1
                                                                                                                                                                                                                                   Query Match 26.1%;
Best Local Similarity 93.3%;
Matches 126; Conservative
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Publication No. US20030037355A1
                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                153 TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT
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mes 126; Conservative
ATCACAAAGGAATCT
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93.3%;
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Pred. No. 1.8e-26;
0; Mismatches 9
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Pred. No. 1.8e-26;
0; Mismatches 9
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; Patent No. US20020004943A1
; GENERAL INFORMATION:
; APPLICAME: ACINOMICS, LLC.
TITLE OF INVENTION: IDENTIFICATION AND CHARACTERIZATION OF A PAGODA PHENOTYPE IN PLAN
FILE REFERENCE: PAGODA
FILE REFERENCE:
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Database
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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462
24791104 seqs, 12571243825 residues
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/cgn2_6/ptodata//pna/US086_COMB.seq:*
/cgn2_6/ptodata//pna/US086_COMB.seq:*
/cgn2_6/ptodata//pna/US089_COMB.seq:*
/cgn2_6/ptodata//pna/US089_COMB.seq:*
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/cgn2_6/ptodata//pna/US090_COMB.seq:*
/cgn2_6/ptodata//pna/US090_COMB.seq:*
/cgn2_6/ptodata//pna/US092_COMB.seq:*
/cgn2_6/ptodata//pna/US093_COMB.seq:*
/cgn2_6/ptodata//pna/US093E_COMB.seq:*
/cgn2_6/ptodata//pna/US095E_COMB.seq:*
/cgn2_6/ptodata//pna/US095E_COMB.seq:*
/cgn2_6/ptodata//pna/US095E_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US084_COMB.seq:*
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44: /cgn2_6/ptodata/1/pna/US6001_COMB.seq:
46: /cgn2_6/ptodata/1/pna/US6001_COMB.seq:
47: /cgn2_6/ptodata/1/pna/US6003_COMB.seq:
48: /cgn2_6/ptodata/1/pna/US6003_COMB.seq:
49: /cgn2_6/ptodata/1/pna/US6003_COMB.seq:
50: /cgn2_6/ptodata/1/pna/US6006_COMB.seq:
51: /cgn2_6/ptodata/1/pna/US6006_COMB.seq:
52: /cgn2_6/ptodata/1/pna/US6006_COMB.seq:
53: /cgn2_6/ptodata/1/pna/US6006_COMB.seq:
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54: /cgn2_6/ptodata/1/pna/US60012_COMB.seq:
55: /cgn2_6/ptodata/1/pna/US6011_COMB.seq:
56: /cgn2_6/ptodata/1/pna/US6012_COMB.seq:
58: /cgn2_6/ptodata/1/pna/US6012_COMB.seq:
59: /cgn2_6/ptodata/1/pna/US6015_COMB.seq:
60: /cgn2_6/ptodata/1/pna/US6015_COMB.seq:
61: /cgn2_6/ptodata/1/pna/US6015_COMB.seq:
62: /cgn2_6/ptodata/1/pna/US6015_COMB.seq:
64: /cgn2_6/ptodata/1/pna/US6017_COMB.seq:
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64: /cgn2_6/ptodata/1/pna/US6017_COMB.seq:
65: /cgn2_6/ptodata/1/pna/US6017_COMB.seq:
66: /cgn2_6/ptodata/1/pna/US6017_COMB.seq:
70: /cgn2_6/ptodata/1/pna/US6013_COMB.seq:
71: /cgn2_6/ptodata/1/pna/US6012_COMB.seq:
73: /cgn2_6/ptodata/1/pna/US6023_COMB.seq:
74: /cgn2_6/ptodata/1/pna/US6023_COMB.seq:
75: /cgn2_6/ptodata/1/pna/US6023_COMB.seq:
76: /cgn2_6/ptodata/1/pna/US6023_COMB.seq:
77: /cgn2_6/ptodata/1/pna/US6031_COMB.seq:
78: /cgn2_6/ptodata/1/pna/US6031_COMB.seq:
79: /cgn2_6/ptodata/1/p
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

1 3 3 4 4 4 4 5 5 6 6 7 7 7 7 10 11 11 11 11 11 11 11 11 11 11 11 11	Result
373.62 354.62 314 262 240.22 212.4 212.4 210.4 210.6 1120.6 1120.6 1120.6 1120.6 1120.6 1120.6	Score
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Sequence 20, Appl Sequence 24, Appl Sequence 19, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 25, Appl Sequence 25, Appli Sequence 6, Appli Sequence 7, Appli Sequence 7, Appli Sequence 1, Appli Sequence 2, Appli Sequence 1, Appli Sequence 4, Appli	Description

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Sequence 20, Application US/09963803

GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

TITLE OF INVENTION: Chimeric expression promoters originating from the promoter of the promoter 
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FEATURE:
FEATURE:
FEATURE:
NAME/KEY: promoter
LOCATION: (1)..(462)
OTHER INFORMATION:
US-09-963-803-20
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  AACAATGAMAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
                                    AACAATGAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
                                                                            GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC
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PCT-US97-10376-2

PCT-US97-10376-2

EUS-09-202-838-3

6 US-09-202-838-3

6 US-09-202-838-3

6 US-09-202-838-3

PCT-US97-10376-3

PCT-US97-10376-3

0 US-09-673-274A-19

5 US-09-673-274A-19

6 US-09-673-274A-19

6 US-09-673-274A-19

5 US-09-847-057-4

10 US-09-847-057-4

10 US-09-847-057-4

10 US-09-103-190-5

PCT-US97-10376-5

PCT-US97-10376-5

PCT-US97-10376A-9

PCT-US97-10376A-9
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Pred. No. 4.4e-123;
; Mismatches 0;
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; NAME/KEY: promoter
; LOCATION: (1)..(600)
; OTHER INFORMATION:
US-09-963-803-22
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US-09-963-803-22
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION UNMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 22
                                                                                                                                                                                                                                                                                                                                                                                        Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: promoter MPr1165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 600
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nilarity 95.3%;
Conservative (
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Pred. No. 2.1e-97;
0; Mismatches 19;
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RESULT 4
US-09-963-803-19
· Sequence 19, Application US/09963803
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US-09-963-803-24
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SEMERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT FILING DATE: 2001-09-26

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR APPLICATION NUMBER: PG 1500/00370

PRIOR APPLICATION NUMBER: PG 1500/00370

PRIOR APPLICATION NUMBER: PG 1500/00370

PRIOR FILING DATE: 2000-10-05

RUMBER OF SEQ ID NOS: 39

CONTENDED TO THE PRIOR PRIOR
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Matches
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SEQ ID NO 24
LENGTH: 541
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; Sequence 23, Application US/09963803
; GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
; CURRENT APPLICATION NUMBER: US/09/963,803
; CURRENT FILING DATE: 2001-09-26
; PRIOR APPLICATION NUMBER: FR 99/03925
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
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LENGTH: 393
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local (
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
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Pred. No. 3.5e-80;
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Sequence 21, Application US/09963803

GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
ITITLE OF INVENTION: Chimeric expression promoters originating from community of inventions of the promoters originating from community of the promoters originating from community of the prince of inventions of the prince of inventions of the prince of invention of the prince of 
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PRIOR APPLICATION NUMBER: PCT IB
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 23
LENGTH: 604
TYPE: DNA
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Best Local Similarity
Matches 375; Conserv
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NAME/KEY: promoter
LOCATION: (1)..(604)
OTHER INFORMATION:
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OTHER INFORMATION: promoter MPr1167
LENGTH:
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Pred. No. 4.9e-65;
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PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 3
LENGTH: 317
TYPE: DNA
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; LOCATION: (1)..(392)
; OTHER INFORMATION:
US-09-963-803-21
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us-09-963-803-3
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Best Local Similarity
Matches 349; Conserv
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
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CURRENT FILING DATE: 2001-09-26
               NAME/KEY: promoter LOCATION: (1)..(317) OTHER INFORMATION:
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                                                                                 OTHER INFORMATION:
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Pred. No. 8.
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR PRIOR APPLICATION NUMBER: PCT 1B00/00370
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-10-05
SOFTMARE: PatentIn version 3.1
SEQ ID NO 25
LENGTH: 472
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Best Local S
Matches 213
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Best Local (
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ORGANISM: Artificial Sequence
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nes 213; Conserv
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                                                                  GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197
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  GGTTCCCAAGCTTTATTT 416
                                                   GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACCCGGTATGCT 398
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Pred. No. 9.6e-51;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         from commelina yellow
                                                                                                                                                                                                                                                             Gaps
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                                                                             317
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                                                                                                      338
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US-09-963-803-4
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LENGTH: 348
                                                              SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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            ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: promoter MPr1146
                                      TYPE: DNA
                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                    172
                                                                                                                                                                                                                                                                                                                  231
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Sequence 4, Application US/09963803
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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                                 APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yell
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: POT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER: OF SEQ. ID NOS: 39
CORPENENCE: DESCRIPTION OF SEQ. ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: promoter MPr1117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   200 GGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 ACTAGTATCCGCCGTCATCAATGACATCACAGCACTGAGGGAGATGAATAGCCTAGCCA 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TAAGGTCGCTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGC
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99.18;
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; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(301)
; OTHER INFORMATION:
US-09-963-803-7
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                                                      RESULT 12
US-09-963-803-6
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PR 99/03925
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-10-05
SOFTWARE: Patentin version 3.1
SEQ ID NO 7
LENGTH: 301
                Sequence 6, Application US/09963803 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 7, Application US/09963803 GENERAL INFORMATION:
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Best Local Similarity 98.7
Matches 154; Conservative
                                                                                                                                                                                                                                                                                                                                                                       Matches 153;
                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: promoter MPr1154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                   138 GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACGACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                            78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                        63 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 122
                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                     GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
                                                                                                                                   GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
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                                                                                                                                                                                                                                                                                                                                                                       Conservative
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98.7%;
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Pred. No. 5.5e-30;
0; Mismatches 2;
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pred. No. 2e-33;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                         Length 301;
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PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEO ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 6
LENGTH: 398
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Best Local Similarity
                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                           APPLICATION NUMBER: US 6
FILING DATE: 20-JUN-1996
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello TITLE OF INVENTION: virus and cassava vein mosaic virus
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                    CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                    ANTI-SENSE:
                                                                                     HYPOTHETICAL:
                                                                                                  TOPOLOGY: 11
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                198 GCGGAAAGTAACCTTATGCATTTGTAATTTTGGTTAC 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    220
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                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US97/10376 FILING DATE: 20-JUN-1997
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                                                                                                                                                                           LENGTH:
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26.1%;
93.3%;
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Pred. No. 6.1e-30;
0; Mismatches 2;
Score 120.6; DB 1;
Pred. No. 4.4e-24;
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                Length 392
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Best Local Similarity

Matches

126;

Conservative

0,

Mismatches

Indels

0;

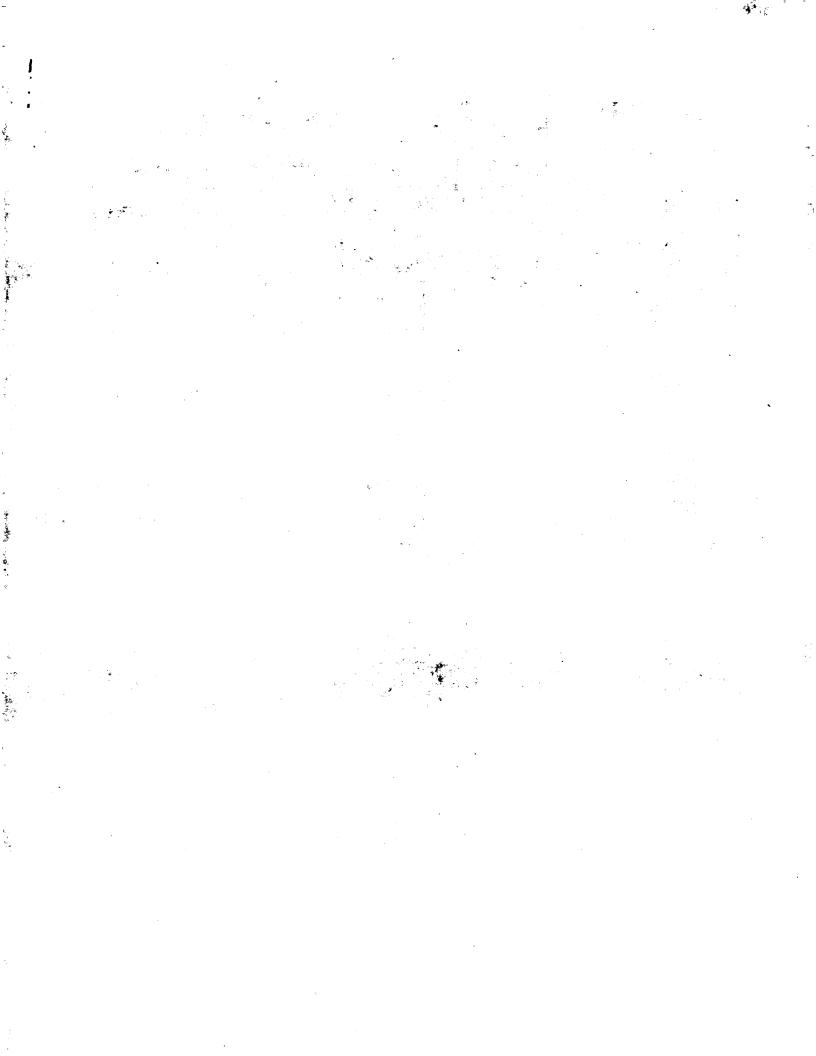
Gaps

0;

GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT 152

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PCT-US97-10376A-1
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PCT-US97-10376A-1
                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/020,129
FILING DATE: 20-JUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US97/103: FILING DATE: 20-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: CASSAVA VEIN MOSAIC VIRUS PROMOTERS AND TITLE OF INVENTION: USES THEREOF NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Beachy, Roger N. APPLICANT: Fauguet, Claude
                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 784-2937
                                                                                                                                                                                                                HYPOTHETICAL:
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                                               153 TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT 212
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CITY: La Jolla
STATE: Californ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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20-JUN-1997
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93.3%;
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Pred. No. 4.4e-24;
0; Mismatches 9
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Search completed: April 14, 2003, 12:06:31 Job time : 1865.88 secs
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; TYPE: DNA
; ORGANISM: Cassava vein mosaic virus
US-09-202-838-1
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US-09-202-838-1
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SEQ ID NO 1
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CURRENT FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/US97/10376
PRIOR FILING DATE: 1997-06-20
PRIOR APPLICATION NUMBER: 60/020,129
PRIOR FILING DATE: 1996-06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Beachy, Roger N.
APPLICANT: Fauquet, Claude
TITLE OF INVENTION: CASSAVA VEIN MOSAIC VIRUS PROMOTERS AND USES THEREOF
FILE REFERENCE: MYCOO11S
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nes 126; Conserv
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de Kochko, Alexandre
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93.3%;
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                 Pending_Patents_NA_New: *
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//cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
//cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
//cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
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     DB
      US-09-765-558-1
US-09-673-274B-20
US-09-673-274B-20
US-10-380-935-30
US-10-380-935-37
US-10-380-935-37
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US-10-160-764-90
US-10-198-478-13
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US-10-198-478-14
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Sequence 1, Appli
Sequence 20, Appl
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Sequence 30, Appl
Sequence 37, Appl
Sequence 147, Appl
Sequence 1, Appli
Sequence 18, Appl
Sequence 18, Appl
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Sequence 40, Appl
Sequence 4, Appli
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Si 6;	1, Applicati INFORMATION: NT: The SCCIE NT: Barbas, NT: Barbas, NT: Guan, Xu NT: Clamia, FINVENTION: FINVENTION: FILING DATE: PLICATION NU PILICATION NU PILING DATE: 2 POF SEQ ID NOS E: FastSEQ fo O 1 DNA SM: Artificia E: 532 DNA SM: Artificia E: INFORMATION: INFORMATION:	4 6 6 6 6 6 6 7 7 7 7 7 7 7 7 7 7 7 7 7
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imilarity 93.3%; Pred, No. 1.4e-25; Conservative 0; Mismatches 9; Indels 0; ATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGA	cation US/09765555 ON: cripps Research It as, Carlos e, Justin , Xueni iia, Bipin ON: Methods and co ON: expression ir 27801-20014.20 110N NUMBER: US/09 ATE: 2001-01-19 N NUMBER: US 60/17 E: 2000-07-21 NUMBER: US 60/17 NOS: 75 NOS: 75 NOS: 75 NOS: 75 O for Windows Vers Con: Promoter CsVM	\$0715 \$0715 206 382 2267 382 382 382 382 382 382 393 4182 4182 4182 5449 5449 55449
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Pred. No. 1.4 Pred. No. 1.4 ; Mismatches ; Mismatches ; Mismatches ; GGACAACAATGAA [CGACAACAATGAA CGACAACAATGAA TGTAAGGTGGAAA TGTAAGGTGGAAA	Institute Institute composit: in plants in plants),897 /177,468 arsion 4.0	H-11-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-
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RESULT 3
US-09-673-274B-20
; Sequence 20, Application US/09673274B
; GENERAL INFORMATION:
                                                                                              APPLICANT: HOFFman, Jules
TITLE OF INVENTION: GENE CODING FOR HELIOMICINE, AND
FILE REFERENCE: A33595-PCT-USA (0726667.0166)
CURRENT APPLICATION NUMBER: US/09/673,274B
CURRENT FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: PCT/FR99/00843
PRIOR FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: PCT/FR99/00843
PRIOR FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: FR 98 04933
PRIOR FILING DATE: 1998-04-15
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TITLE OF INVENTION: GENE CODING FOR HELIOMICINE, AND USE THEREOF
FILE REFERENCE: A33595-PCT-USA (0726667.0166)
CURRENT APPLICATION NUMBER: US/09/673,274B
CURRENT FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: PCT/FR99/00843
PRIOR FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: FR 98 04933
PRIOR APPLICATION NUMBER: FR 98 04933
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 48
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 19
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Best Local Similarity
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                                                                                       NUMBER OF SEQ ID NOS: 48
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TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 838
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_structure LOCATION: (533)...(568)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: promoter LOCATION: (7)...(532) FEATURE:
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                        1036
                                                                 FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                 Bulet, Phillipe
Brookhart, Gary
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93.3%;
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Pred. No. 1.6e-25;
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CURRENT APPLICATION NUMBER: US/10/380,935
CURRENT FILING DATE: 2003-03-19
PRIOR APPLICATION NUMBER: GB0023911.1
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: GB0027693.1
PRIOR FILING DATE: 2000-11-13
PRIOR FILING DATE: 2000-99-29
NUMBER OF SEQ ID NOS: 70
SOSTWARE: Patentin version 3.1
SEQ ID NO 30
LENGTH: 309
TYPE: DNA
US-10-380-935-37

Sequence 37, Application US/10380935

GENERAL INFORMATION:

APPLICANT: WARNEE, Simon, Anthony, Applicant: HAWKES, Timothy, Robert
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; LOCATION: (767)...(1030)
US-09-673-274B-20
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Best Local Similarity
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                                                                                                                                                                                                                                                                       Query Match
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APPLICANT: HAWKES, Timothy, Robert
APPLICANT: ANDREWS, Christopher, John
TITLE OF INVENTION: HERBICIDE RESISTANT PLANTS
FILE REFERENCE: PPD50594/WO
                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Cauliflower Mosaic Virus FEATURE:
                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Enhancer
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NAME/KEY: CDS
LOCATION: (539)...(736)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                        342
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                                                                                                                                                                                                                                                      y Match 8.7%;
Local Similarity 88.0%;
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                                                                                                                                                                                                                                    44;
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93.3%;
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                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                       Length 309;
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Anthony, James

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CURRENT APPLICATION UNMBER: US/10/380,935
CURRENT FILING DATE: 2003-03-19
PRIOR APPLICATION NUMBER: GB0023911.1
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION UMBER: GB0023910.3
PRIOR FILING DATE: 2000-11-13
PRIOR FILING DATE: 2000-09-29
NUMBER GB023910.3
PRIOR FILING DATE: 2000-09-29
UNMBER OF SEQ ID NOS: 70
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 37

                                                                                                                                                                                                                        US-60-434-166-147
US-60-434-166-147;
Sequence 147, Application US/60434166;
GENERAL INFORMATION:
APPLICANT: Mendel Biotechnology
Creelman, Robert A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 10
; LENGTH: 835
; TYPE: DNA
; ORGANISM: Cauliflower mosaic virus
US-10-391-414-10
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: SHINOZAKI, Kazuko
APPLICANT: KATSURA, Koji
APPLICANT: ITO, YUSUKe
TITLE OF INVENTION: Stress Induced Promoter Derived From Rice
FILE REFERENCE: 382.1041
CURRENT APPLICATION NUMBER: US/10/391,414
CURRENT FILING DATE: 2003-03-18
                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: JP 2002-377316 PRIOR FILING DATE: 2002-12-26 NUMBER OF SEQ ID NOS: 11
                                                                        APPLICANT:
                                                                                                          APPLICANT:
                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ANDREWS, Christopher, John
TITLE OF INVENTION: HERBICIDE RESISTANT PLANTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                              Haake, Volker
Ratcliffe, Oliver
                          Jiang, Cai-Zhong
Reuber, T. Lynne
                                                                                                      Heard, Jacqueline E. Adam, Luc J.
Riechmann, Jose Luis
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                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/302,267
CURRENT FILING DATE: 2002-11-22
PRIOR APPLICATION NUMBER: US/09/506,720
PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/120,880
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: 60/121,037
PRIOR FILING DATE: 1999-02-22
PRIOR FILING DATE: 1999-02-22
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SEQ ID NO 147
LENGTH: 1683
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                                                                                                                              SEQ ID NO 1
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Best Local Similarity 88.0%;
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APPLICANT: CreeLman, Robert
TITLE OF INVENTION: PLANT GENE SEQUENCES II
FILE REFERENCE: MBI-0007
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                                                                                                                                                SOFTWARE: PatentIn
                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/161,143 PRIOR FILING DATE: 1999-10-22
                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 60/144,153
PRIOR FILING DATE: 1999-07-15
                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/135,134
                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/124,278
PRIOR FILING DATE: 1999-03-11
PRIOR APPLICATION NUMBER: 60/129,450
PRIOR FILING DATE: 1999-04-15
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CURRENT FILING DATE: 2002-12-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES IN PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                         PRIOR FILING DATE:
                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/162,656
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NAME/KEY: CDS
LOCATION: (144)..(2024)
                                         FEATURE:
                                                             TYPE: DNA ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                        2361
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Zhang, James
Yu, Guo-Liang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jiang, Cai-Zhong
Samaha, Raymond
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Heard, Jacqueline
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Pineda, Omaira
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Adam, Luc
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                                                                                                                                                                                         1999-11-01
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RESULT 11
US-10-130-150-15
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Sequence 15, Application US/10130150 GENERAL INFORMATION:
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                                                                                                                                                                   Query Match
Best Local Similarity
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Best Local :
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TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THEREOF
FILE REFERENCE: 02/23831
CURRENT APPLICATION NUMBER: US/10/130,150
CURRENT FILING DATE: 2000-11-15
NUMBER OF SEQ ID NOS: 24
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TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THEREOF
FILE REFERENCE: 02/23831
CURRENT APPLICATION NUMBER: US/10/130,150
CURRENT FILING DATE: 2000-11-15
NUMBER OF SEQ ID NOS: 24
                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                            FEATURE:
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                                                                                                                                                                                                                                                                                                        LENGTH: 3288
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                                                                                       258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
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les 44; Conservative
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Pred. No. 0.14;
0; Mismatches 6;
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Pred. No. 0.13;
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Pred. No. 0.14;
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US-10-160-764-40
US-10-60-764-40
Sequence 40, Application US/10160764
GENERAL INFORMATION:
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; OTHER INFORMATION:
US-10-160-764-40
                                                                       RESULT 13
US-10-160-764-4
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                                   Sequence 4, Applicat GENERAL INFORMATION:
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SEQ ID NO 40
LENGTH: 5247
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Best Local Similarity
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TITLE OF INVENTION: Compositions and Methods
TITLE OF INVENTION: in Plants
FILE REFERENCE: 22542-008
CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
PRIOR FILING DATE: 2001-05-31
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TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THER
EILE REFERENCE: 02/23831
CURRENT APPLICATION NUMBER: US/10/130,150
CURRENT FILING DATE: 2000-11-15
                  APPLICANT: Huang,
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 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 90
                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                 3209 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 3258
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les 44; Conserv
                                                    Application US/10160764
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Chalifoux, Maryse
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                                                                                                                                                                                                                      Conservative
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                    Yafan
                                                                                                                                                                                                                                                                                                           Description of Artificial Sequence: Plasmid pBI121-35S-AtFTA
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APPLICANT: Gilley, TITLE OF INVENTION:

Angela Patricia Compositions and

Methods of Increasing Stress Tolerance

APPLICANT:

Wang, Yang Kuzma, Monika Maria

TITLE OF INVENTION: in Plants

FILE REFERENCE:

22542-008

CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR FILING DATE: 2001-10-22
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENGTH: 5250
TYPE: DNA
ORGANISM: Artificial Sequence

Query Match

Length 5250;

OTHER INFORMATION: Description of Artificial Sequence: Plasmid OTHER INFORMATION: pBI121-35S-anti-AtFTA sequence

FEATURE:

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RESULT 15
US-10-160-764-52
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                                                    Sequence 52, Application US/10160764 GENERAL INFORMATION:
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SEQ ID NO 48
LENGTH: 5511
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity 88.0%;
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CURRENT FILLING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR FILING DATE: 2001-10-22
NUMBER OF SEQ ID NOS: 90
            APPLICANT: Huang, Yafan APPLICANT: Chalifoux, Maryse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gilley, Angela Patricia
TITLE OF INVENTION: Compositions and Methods of Increasing Stress Tolerance
TITLE OF INVENTION: in Plants
FILE REFERENCE: 22542-008
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APPLICANT: Chalif
APPLICANT: Wang,
APPLICANT:
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APPLICANT: Gilley, Angela Patr
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                                                                                                                                                                    3209 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 3258
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Local Similarity 88.0%;
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Pred. No. 0.16;
0; Mismatches 6,
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   Search completed:
Job time: 377.79
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                                                                                                                                                         Matches
                                                                                                                                                                         Query Match 8.7%;
Best Local Similarity 88.0%;
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APPLICANT: Gilley, Angela Patricia
TITLE OF INVENTION: Compositions and Methods of Increasing Stress Tolerance
TITLE OF INVENTION: in Plants
FILE REFERENCE: 22542-008
                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/348,909 PRIOR FILING DATE: 2001-10-22 NUMBER OF SEQ ID NOS: 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/160,764 CURRENT FILING DATE: 2002-05-31
                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                     OTHER INFORMATION:
OTHER INFORMATION:
                                                                               3209 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 3258
                                                                                                   258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                                                                                                                                                                                                                                                                                                      PatentIn Ver.
                                                                                                                                                         Conservative
April 14, 2003, 12:27:40 secs
                                                                                                                                                                                                                                                     Description of Artificial Sequence: Plasmid pBI121-35S-AtFTB
                                                                                                                                                                                                                                                                                                                                                                                            2.1
                                                                                                                                                       Score 40.4; DB Pred. No. 0.17; 0; Mismatches
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                                                                                                                                                                                           Length 5715;
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Maximum DB seq length: 2000000000
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Maximum Match 100%
Listing first 45 summaries
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                                                                                      Score
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Match
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BH746858 SALK_0036
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AL529655 AL529655
AL063921 Drosophil
BM161725 EST564248
BH751161 SALK_0494
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SALK_0089	9	8	9	245	9	360	337	246	26H	8	250	86	90	031	26 H	26F	046	SALK_0460	337	217	20	26E	25B	242	86B	217	$\frac{\omega}{\omega}$	229	35	245	219	26E	24.5	516	245	800	240	047	

ALIGNMENTS

	TITLE JOURNAL COMMENT	ORGANISM REFERENCE AUTHORS	ACCESSION ACCESSION VERSION KEYWORDS SOURCE
Salk Institute Genomic Analysis Laboratory (SIGnAL) The Salk Institute for Biological Studies 10010 N. Torrey Pines Road, La Jolla, CA 92037, USA Tel: 858 453 4100 x1752 Fax: 858 558 6379 Email: ecker@salk.edu This is single pass sequence recovered from the left border of	, Zimmerman, J. and Ecker, J.R. A Sequence-Indexed Library of Insertion Mutations in the Arabidopsis Genome Unpublished (2001) Contact: Joseph R. Ecker	Arabidopsis thaliana Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis. (bases 1 to 224) Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Prednis,L., Shinn,P.	BH746858 224 bp DNA linear GSS 27-FEB-2002 SALK_003694.51.40.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_003694.51.40.x, DNA sequence. BH746858 BH746858.1 GI:18959973 GSS. thale Cress.

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SOURCE
ORGANISM
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BM162517
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      297 TGCCACGACTA 307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Jane Carlton
Parasite Genomics Group
The Institute for Genomic Research
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Plasmodium yoelii yoelii
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BM162517.1 GI:17308198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Plasmodium yoelii EST project at TIGR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
1 (bases 1 to 561)
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                                                                                                                                                                                                                                                                                                                                                                       For clone info, please contact the Malaria Research Reagent Resource Center, ATCC
                                                                                                                                                                                                                                                                                                                                                                                                              Email: carlton@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 Medical Center Drive,
1: 301-530-9319
                                                                                                                                                                                                                                                                                                                              primer: ADF.
                                                                                                                                                                                                                                                                                                                                                                                                                                    301-838-0208
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/note="PCR was performed on Arabidopsis thallana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 68 c 54 g 50 t
/dev_stage="Asexual blood stages"
/lab_host="E. coli XL-1 Blue"
/note="Vector: pAD-GAL4; At 20-25% parasitemia, blood collected from BALB/CByJ mice infected with Py17XL parasites, and leukocytes removed by passage over microcrystalline cellulose collumns. Total RNA was isolated using the guanidinium isothlocyanate method, mRNA isolated using oligo(dT)-cellulose chromatography
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/db_xref="taxon:3702"
/clone="SALK_003694.51.40.x"
                                                                                                                                                                                                                                           /organism="Plasmodium yoelii yoelii"
/strain="17XL"
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                                                                                                                                                                            /clone_lib="PyBS"
                                                                                                                                                                                               /clone="PYCKT64"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rockville, MD 20850, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               82 TGAAAATAACGCAAAATCGGAATTTATGAAAATGGCCCTAGCAGGTACATAAAATGAATAG 141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Li,W.B., Gruber,C., Jessee,J. and Polayes,D. Full-length cDNA libraries and normalization Unpublished (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 964)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AL529655.1 GI:12793148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Genoscope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prime, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genoscope - Centre National de Sequencage
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       257
/note="Organ: brain; Vector: pCMVSPORT 6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371 Email: fliang@lifetech.com URL: http://fulllength.invitrogen.com"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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segref@genoscope.cns.fr, 1
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                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="CSODD005YL16"
                                                                                                                                                                                                                                                            /tissue_type="neuroblastoma cells"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                          /sex="male"
                                                                                                                                                                                                                                                                                                                                 /clone_lib="LTI_NFL001_NBC4"
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Query Match 9.0%;
Best Local Similarity 17.3%;
Matches 39; Conservative 10
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Best Local Similarity
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                                                                                                                                                                                                                          Determination of this BAC-end sequence was carried out as part of a collaboration with the Berkeley Drosophila Genome Project (BDGP). The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial EcoRI digestion of Drosophila DNA provided by the BDGP from the isogenic Strain y2; cn bw sp, the same strain used for the BDGP's Pl and EST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be cocation/Qualifiers
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fi
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AL063921.1 GI:4941778
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BACR08K10 of RPCI-98 library from
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                                                                                            /clone="BACR08K10"
/clone_lib="RPCI-98"
/note="end : TET3"
a 64 c 131 g
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/db_xref="taxon:7227"
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107; Mismatches
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                                    Length 1101;
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Carlton,J.M., Daly,T.M., Long,C.A., Bergman,L.W., Vaidya,A.B.,
Fraser,C.M. and Carucci,D.J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: carlton@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Parasite Genomics Group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plasmodium yoelii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BM161725.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reagent Resource Center, ATCC
                            99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     301-838-0208
                            ρ
                                                                                                                  5-methyl dCTP. After second strand synthesis, uneven termini were treated with Pfu DNA polymerase and EcoRI adaptors ligated to the blunt ends. The sample was cleaved with XhoI and separated on a Sephacryl S-500 column. Size-fractionated cDNA was precipitated and ligated to HybrizAP arms directionally using EcoRI-XhoI cleaved arms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jane Carlton
                            isolated."
25 c
                                                                           excised
                                                                                                                                                                                                                                                                                             isolated using the guanidinium isothiocyanate method, amRNA isolated using oligo(dT)-cellulose chromatography First strand cDNA synthesis was completed using a 50-ba
                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="Asexual blood stages"
/lab_host="E. coli XL-1 Blue"
/note="Vector: pAb-GAL4; At 20-25% parasitemia, blood was
collected from BALB/CByJ mice infected with Py17XL
parasites, and leukocytes removed by passage over
microcrystalline cellulose collumns, Total RNA was
                                                                                                After packaging,
                                                                                                                                                                                                                                                                         primer and reverse transcriptase in the presence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:73239"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /strain="17XL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="PyBS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="PYCKI31"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Plasmodium yoelii yoelii"
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                                                                        from the HybriZAP vector and
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                                                                                                  the phagemid vector (pAD-GAL4) was
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                           GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGCGCAAGCAATCATATTATATTATGCCTTATTATTTATG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2001)
Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
The Salk Institute For Biological Studies
The Salk Institute For Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Predni, Zimmerman,J. and Ecker,J.R.

A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence.
BH751161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10010 N. Torrey Pines Road,
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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GSS.
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                                                                                                                                                                                                                                           /db_xref="taxon:3702"
/db_xref="taxon:3702"
/clone="$ALK_049464.49.75.x"
/clone="$ALK_049464.49.x"
/clone="$ALK_049
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                                                                                                                                                          Score 40.4; D
Pred. No. 2.8;
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Prednis, L., Shinn, I
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                        AUTHORS
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Alonso,J.M., Leisse,T.J.,
,C., Jeske,A., Karnes,M.,
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GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA
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                                                                                                                                                                                                                                           H619283 153 bp DNA' linear GS SALK_040762 Arabidopsis thaliana TDNA insertion lines thaliana genomic clone SALK_040762, DNA sequence. BH619283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 888 453 4100 x1752
Fax: 858 558 6379
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                                                                     Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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A Sequence-Indexed Library of Insertion Mutations in the
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 36 c 30 g 35 t
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/db_xref="taxon:3702"
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/strain="Columbia 0"
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88.0%;
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RESULT 9
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BH747013 153 bp DNA linear GSS 27-FEI SALK_008070.43.05.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_008070.43.05.x, DNA
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Arabidopsis Genome
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 153)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadr
                                                                                                                                                                                                                                                                                                              Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Prednis, L., Zimmerman, J. and Ecker, J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                              Class: TDNA tagged
                                                                                                                            Email: ecker@salk.edu
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 34 c 36 g 47 t
                        Location/Qualifiers
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/organism="Arabidopsis thaliana"
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Library of Insertion Mutations in the
                                                                                                      sequence recovered
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., G., C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Zimmerman,J. and Ecker,J.R.

A Sequence-Indexed Library of Insertion Mutations in the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arabidopsis Genome 
Unpublished (2001)
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Alonso, J.M., Leisse, T.J.,
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BH748291
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Tel: 858 453 4100 x1752
Fax: 858 558 6379
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/clone=lib="Arabidopsis thaliana TDNA insertion lines"
/clone=PPCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 37 c 33 g 36 t
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 35 c 37 g 49 t
                                                                                                                                                                                              /strain="Columbia 0"
/db_xref="taxon:3702"
/clone="SALK_045100.51.10.x"
                                                                                                                                                                                                                                                               /organism="Arabidopsis thaliana
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Query Match Best Local Similarity

8.7%;

Score 40.4; DB 17; Pred. No. 2.9;

Length 158;

Conservative

0;

6,

Indels

0

Gaps

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RESULT 12
BH746616
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Best Local :
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                                                                                                                    BH746616

175 bp DNA IIIIeaa CONSALK_045268.54.50.x Arabidopsis thaliana TDNA insertion SALK_045268.54.50.x,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R.,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
Zinmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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                                                                                                sequence.
BH746616
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1 (bases 1 to 165)
                                                                              BH746616.1
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                                           thale cress
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      thale cress
                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="SALK_016522.55.50.x"
/clone_lib="Arabidopsis thaliana Innes"
/clone_lib="Arabidopsis thaliana lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 43 c 39 g 38 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Arabidopsis thaliana"
/strain="Columbia 0"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                              GI:18959731
                                                                                                                                                                                                                                                                                                                                               88.7%;
                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                             Score 40.4;
Pred. No. 2
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Prednis, L., Shinn,
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AUTHORS
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JOURNAL
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Best Local :
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                                                                                                                                                                                                                                                      Walbot, V
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Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Prednis, L., Zimmerman, J. and Ecker, J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                               M85 California Ave, Palo Alto, CA 94304, Tel: 650 723 2227 Fax: 650 725 8221 Email: walbot@stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis 1 (bases 1 to 175)
Class: transposon-tagged.
Location/Qualifiers
                                                                                                                                                                                                           Department of Biological Sciences Stanford University
                                                                                                                                                                                                                                                         Contact: Walbot V
                                                                                                                                                                                                                                                                                  Unpublished (2001
                                                                                                                                                                                                                                                                                                                                                        clade; Panicoideae; Andropogoneae;
1 (bases 1 to 177)
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                                              Plate: 1008026 row:
                                                                   Very probable ligation site found so sequence was trimmed Post-ligation sequence submitted separately.
                                                                                                                                                                                                                                                                                                          Maize genomic sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /OLONE="SAKE 045268.54.50.x"
/Clone="Ib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 43 c 40 g 38 t
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/db_xref="taxon:3702"
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RescueMu Grid
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Pred. No. 3;
0; Mismatches
                                                                                                                                                                                                                                                                                                        found using
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Prednis, L., Shinn,
                                                                                                                                                                                                                                                                                                            RescueMu transposon
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JOURNAL COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AUTHORS
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Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BH752801 190 bp DNA linear GSS SALK_019366.54.25.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK_019366.54.25.x,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H.,,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GSS
                                                                                                                                                                                                    Class: TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; euroside; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                        This is single pass
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence-Indexed Library
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Zimmerman J. and Ecker,J.R.
Zimmerman J. and Ecker,J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 190)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site_1: BamHI; Site_2: BglII; RescueMu is a 4.9 kb, modified maize Mu transposon designed to allow plasmid rescue from total genomic DNA. Mu elements insert preferentially into transcription units. For more information on RescueMu, go to the web site www.zmdb.iastate.edu and follow the links for 'RescueMu.' Grid I was grown at Berkeley in 2001. DNA was extracted from leaf punches, double djegsted using BamHI and BglII, and ligated to form circular plasmids. DH10B cells were transformed and then screened on LB plates with ampicillin."
                                                                                                                                                                                                                                                                 ecker@salk.edu
              /Clone="SALK_019366.54.25.x"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /cultivar="mixed background W23/A188/B73"
/db_xref="taxon:4577"
/clone_llb="1008 - RescueMu Grid I"
                                                                                                                      /strain-"Columbia
                                                                                                                        /organism="Arabidopsis thaliana"
/strain="Columbia 0"
                                                                                                                                                                             Location/Qualifiers
                                                                                                    /db_xref="taxon:3702"
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88.0%;
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    The resultant fragment for each
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84 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 133
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Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
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A Sequence-Indexed Library of Insertion Mutations in
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                                                                                                                                                                                          /Clone="SALK_045097.54.25.x"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
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SUMMARIES

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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 57
AX093052
AX093052.1
1 (bases 1 to 15077)
Gruber, V. and Comeau, D.
Synthetic vectors, transgenic plants
for obtaining them
Patent: WO 0118192-A 57 15-MAR-2001;
MERISTEM THERAPPUTICS (FR)
Location/Qualifiers
                                                                 artificial sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1. .9285
/organism="synthetic c
/db_xref="taxon:32630"
                                                                            construct
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/ 100.0%; Pi
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Pred. No.
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587; Conserv
        Rance.I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 20 05-0CT-2000;
                                              synthetic construct artificial sequences.

1 (bases 1 to 462)
                                                                                                        AX036754
Sequence 20 from
AX036754
  MERISTEM THERAPEUTICS
                                                                           synthetic construct.
                                                                                              AX036754.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="synthetic construct"
/db_xref="taxon:32630"
/note="pMRT1342"
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Pred. No. 2.2e-150;
 (FR) ; RANCE
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Sequence 23
AX036757
                                                                                                                                1 (bases 1 to 604)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus patent: WO 0058485-A 23 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
                                                                                                                                                                                                artificial sequences.
                                                                                                                                                                                                          synthetic construct
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/note="promoter MPr1163"
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/db_xref="taxon:32630"
                                                                      /note="promoter MPr1167"
                                                                                                             Location/Qualifiers
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Pred. No. 6.1e-75;
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Pred. No. 8.9e-92;
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              AAGCTTGCATGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA
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                                              316;
                                                                                                                                                                                                                       Sequence 19
AX036753
AX036753.1
                                                                                                                                     Rance, I., Theisen, M. and Gruber, V.
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
patent: WO 0058485-A 19 05-OCT-2000;
MERISTEM THERAPPUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
                                                                                                                                                                                                synthetic construct. synthetic construct
                                                    Similarity
                                                                                                                                                                                         artificial sequences.
                                                                                                                                                                                                                                              AX036753
                                                                                  128
                                             Conservative
                                                                                               /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1162"
                                                                                                                        Location/Qualifiers
1. .393
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AX036758
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CGGTATGCTGGTTCCCAAGCTTTATTT 485
                        TGATTGATGTGATATCAAGATTGATGT 404
                                                                                                  GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
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AX036758
AX036758.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      synthetic construct. synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               artificial sequences.
1 (bases 1 to 541)
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/db_xref="taxon:32630"
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REFERENCE
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AX036737
LOCUS
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BASE COUNT
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                                                 synthetic construct.

M synthetic construct
artificial sequences.

1 (bases 1 to 317)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating fro
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 3 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR)
; GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                           AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGACTAGT
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Sequence
AX036755
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Sequence 3 from Patent W00058485
AX036737
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1 (bases 1 to 392)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 21 05-OCT-2000;
                                                                                                                                                                                          AX036737.1
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/db_xref="taxon:32630"
/note="promoter MPr1164"
1. .392
  /organism="synthetic construct"
/db_xref="taxon:32630"
                                       Location/Qualifiers
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Pred. No. 1.5e-53;
0; Mismatches 8
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  TGATATCAAGATTGATGT 335
                           GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGTTACCCCGGTATGCT 398
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213; Conser
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Sequence
AX036759
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Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 25 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                         artificial sequences.
1 (bases 1 to 472)
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25 from Patent WO0058485.
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/db_xref="taxon:32630"
/note="promoter MPr1169"
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Pred. No. 1.5e-47;
0; Mismatches 21
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Pred. No. 1.5
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artificial sequences.
1 (bases 1 to 348)
2 1 (bases 1 to 348)
3 Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from commottle virus and cassava vein mosaic virus
patent: WO 0058485A 4 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THI
; GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                      GGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 264
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                                                    Rance,I., Theisen,M. and Gruber,V. Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 5 05-OCT-2000; MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); GRUBER VERONIQUE (FR);
                                                                                                                                                                                      Sequence 5 from Patent WO0058485
                                                                                                                                            synthetic construct. synthetic construct
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Sequence 4 from Patent WO0058485,
AX036738
AX036738.1 GI:11226247
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1 (bases 1 to 371)
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/db_xref="taxon:32630"
/note="promoter MPr1117"
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70 c 78 g 84 t
 /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1146"
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Promoter
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AX036740
Sequence 6
AX036740
AX036740.1
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AX036741
AX036741.1
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Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating fromottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 7 05-0CT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
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 GI:11226249
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Pred. No. 6.5e-28;
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Pred. No. 3.3e-31;
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Matches 153; Conserv
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synthetic construct
artificial sequences.
1 (bases 1 to 515)
Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 2 05-CCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR); GRUBER VERONIQUE (FR)
COCATION/QUALIfiers
1 616
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Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 6 05-0CT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR); GRUBER VERONIQUE (FR)
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ilarity 98.1%;
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/db_xref="taxon:32630"
/note="promoter MPr1147"
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a 80 c 93 g 97 t
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/db_xref="taxon:32630"
/note="Promoter from the intergenic region of Cassava Vein Mosaic Virus of 515 bp in length EMBL U59751"
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Qy 213 ATGCATTTGTAATTT 227
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Db 336 ATCACAAAGGAATCT 350
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Search completed: April 14, 2003, 09:29:40 Job time: 1728.81 secs

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Result
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ALIGNMENTS

RESULT 1 AAA96856 Chimeric - Commelina yellow mottle virus Chimeric - Cassava vein mosaic virus. Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss. WPI; 2000-647238/62 Rance I, 05-OCT-2000. WO200058485-A1 Nucleotide sequence of chimeric expression promoter MPr1165 19-FEB-2001 AAA96856; AAA96856 standard; DNA; 600 29-MAR-1999; 29-MAR-2000; 2000WO-IB00370 (MERI-) MERISTEM THERAPEUTICS Gruber V, (first entry) 99FR-0003925 ВP

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ACGACTAGTGATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATG
                                                                                                                                                                                                                                 AAGCTTGCATGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5
                                                                                                                                                                                                                                                                                                                                                  from intergenic regions. The chimeric ducing transgenic plants.
                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                            462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expression promoter for transgenic plant production, comprises from promoter comprising vascular expression region replaced uence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    •
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Commelina
Cassava ve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence of chimeric expression
                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                            BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99FR-0003925
                                                                                                                                                                                                                                                                                                                            148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     91pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vein
                                                                                                                                                                                                                                                                                         62.3%;
95.3%;
                                                                                                                                                                                                                                                                                                                           A; 87 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            yellow mottle virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mosaic
                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                             Score 373.6;
Pred. No. 2e-1
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              X
                                                                                                                                                                                                                                                                                                                            111 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   virus
                                                                                                                                                                                                                                                                                                                            116 T; 0 other;
                                                                                                                                                                                                                                                                                          2e-104;
                                                                                                                                                                                                                                                                                                     DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            promoter
                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                            promoters
                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                    Length
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                                                                                                                                         Query Match
Best Local S
Matches 442
                                                                                                                                                                                                                         The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                         Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric - Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric expression promoter; plant green tissue expression
                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plant green tissue expression transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA96857;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA96857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleotide sequence of chimeric expression promoter MPrl167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-FEB-2001
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               198
                                        220
                                                                                         160
                                                                                                                 78
                                                                                                                                                      Local Similarity
GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000-647238/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACGCATGCCACGACTAGTGATTGATGTGATATCAAGATTGATGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MERISTEM THERAPEUTICS.
                                                                                                                                                                                            604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Commelina yellow mottle virus
                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                               sequence represents a chimeric promoter of the invention
                                                                                                                                                                                           ₽₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                        87; 91pp;
                                                                                                                                                                                                                    transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99FR-0003925
                                                                                                                                                                                        186 A; 116 C; 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                   52.0%;
84.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Theisen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         604
                                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mosaic virus
                                                                                                                                                                                                                   plants.
                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                     Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plant vascular expression promoter; promoter; Cassava vein mosaic virus
                                                                                                                                          Mismatches
                                                                                                                                    e 312.2; L.
No. 1.4e-85;
                                                                                                                                                                                        G; 157 T; 0 other;
                                                                                                                                                              21; Length 604;
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                                                                                                                                         Indels
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RESULT 4
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                                                                                                                                                                                                                                                                 Chimeric expression promoter for transgenic plant production, compris-
sequence from promoter comprising vascular expression region replaced
with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                       WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chimeric - Commelina yellow mottle virus Chimeric - Cassava vein mosaic virus.
                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plant green tissu
transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                       Rance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric expression promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleotide sequence of chimeric expression promoter MPr1162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA96853 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAR-2000; 2000WO-IB00370
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                                                                                                                                                                                                 5
                                                                                                                                                                                               Page 85;
                                                                                                                                                                                                                                                                                                                                                                                                                       Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tissue expression plant; chimera; ss.
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                                                                                                                                                                                                                                                                                                                    comprises
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The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from

from

The present sequence represents a chimeric promoter of the invention

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RESULT 5
AAA96858
AID AAA9
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XX 19-F
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KW Prom
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Best Loc
Matches
                                          Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
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Chimeric - Cassava
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ilarity 94.3%;
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Mismatches
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Chimeric -Chimeric -

Cassava

Commelina yellow mottle virus

vein mosaic virus

transgenic plant;

chimera;

05-OCT-2000

WO200058485-A1

(MERI-) MERISTEM THERAPEUTICS

29-MAR-1999; 29-MAR-2000;

2000WO-IB00370 99FR-0003925

Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus

virus;

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RESULT 6
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                                                                                                                                                                                                                                      19-FEB-2001
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                                                                                                                                                                                                                             chimeric expression promoter; plant vascular expression plant green tissue expression promoter; Cassava vein mo: transgenic plant; chimera; ss.
                                                                                                                                                     Chimeric -
Chimeric -
  29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                                  Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                          Nucleotide sequence of chimeric expression promoter MPr1116
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                                                                                                                                                  Commelina yellow mottle virus. Cassava vein mosaic virus.
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                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transgenic plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 234.2; DB 21;
Pred. No. 9.6e-62;
Pred. No. 9.6e-8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 21;
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                                                                                                                                                                                                                                                        ion promoter;
mosaic virus;
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                 Chimeric -
Chimeric -
                                                                                                                     chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus;
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                                         WO200058485-A1
                                                                                                         transgenic plant;
                                                                                                                                                Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                           Nucleotide sequence of chimeric expression promoter MPr1169.
                                                                                                                                                                                                      19-FEB-2001
                                                                                                                                                                                                                                                            AAA96859 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 317 BP; 107 A; 61 C; 74 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   derived from intergenic regions. The chimeric promoters for producing transgenic plants.
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                                                                   Cassava
                                                                                 Commelina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99FR-0003925
                                                                                                           chimera;
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                                                                 vein
                                                                                                                                                                                                     entry)
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99.5%;
                                                                                 yellow mottle virus
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                                                                 mosaic virus
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ARBSULT 9
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Chimeric
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
                                    plant green
transgenic p
                                                            Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression pro
                                                                                                                            19-FEB-2001
                                                                                                                                                                             AAA96838 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   producing transgenic plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             472 BP;
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 Commelina yellow mottle virus Cassava vein mosaic virus.
                                     plant;
                                                                                                   sequence
                                     tissue expression plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000WO-IB00370
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                                                                                                                                                                             DNA;
                                                                                                  of chimeric
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                                                                                                  expression promoter
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No. 4.9
                                                 Cassava vein mosaic virus;
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                                                              promoter;
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Best Local
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                                  chimeric expression promoter; plant vas
plant green tissue expression promoter;
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                        transgenic
                                                           Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                    Nucleotide
                                                                                                              19-FEB-2001
                                                                                                                                      AAA96839;
                                                                                                                                                               AAA96839 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 348 BP; 116 A; 70 C; 78 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gruber V,
                       plant;
                                                                                   sequence
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                       tissue expression
plant; chimera; ss.
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                                                                                                                                                               ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                       200.4;
No. 2.
                                               vascular expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84 T; 0 other;
                                     Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                     .1e-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                   21;
                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                      MPr1146
                                               promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                          1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          comprises
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                           111
                                                                                                                                                                                                                                                                                                   199
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Chimeric

Commelina yellow mottle virus

Chimeric - Cassava

vein

mosaic virus.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 154
                                                             chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                         Chimeric -
Chimeric -
 WO200058485-A1
                                                                                                    Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                              Nucleotide sequence of chimeric expression
                                                                                                                                                       19-FEB-2001
                                                                                                                                                                                                       AAA96841 standard; DNA; 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-647238/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                                                         190
                                                                                                                                                                                                                                                                                                                                                                       138
                                                                                                                                                                                                                                                                         250
                                                                                                                                                                                                                                                                                                198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present sequence represents a chimeric promoter of the invention
                                                                                                                                                                                                                                                                                                                       GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
                                                                                                                                                                                                                                                                                                                                                GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                      GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             371 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gruber V,
                        Commelina yellow mottle virus Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              122 A; 68 C; 89 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                       25.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Theisen
                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 152.8; DB 21
                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 8.2
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ŧ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 21;
                                                                                                                            promoter MPr1154
                                                                                                                                                                                                                                                                                                                                                                                                                             2;
                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                   371;
                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                        249
                                                                                                                                                                                                                                                                                                                                                197
                                                                                                                                                                                                                                                                                                                                                                                                                           0
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ARASJET 12
ARASJES 40
ID ARASJES XX
AC ARASJES XX
DT 19-FE
XX
Promo
KW Promo
KW Chime
KW Plant
KW trans
XX
Chime
OS Chime
OS Chime
OS Chime
XX
WO200
XX
WO200
PD 05-OC
XX

chimeric expression promoter; plant vaso plant green tissue expression promoter;

intergenic region; Commelina yellow mottle

plant vascular expression

Cassava vein mosaic virus;

promoter;

transgenic plant;

chimera;

SS.

Nucleotide sequence of chimeric expression

promoter MPr1147

19-FEB-2001

(first entry)

AAA96840 standard; DNA; 398

ВP

Chimeric Chimeric

Commelina yellow mottle virus Cassava vein mosaic virus.

WO200058485-A1

. Qy В

198 GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233

GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG

181

1;

GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC

123

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δÃ
                                          멍
                                                                 δÃ
                                                                                                                                                          Query Match
                                                                                                Matches
                                                                                                                                                                                                             The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mossic virus. Especially, the promoters are
                                                                                                                                                          Sequence 301
                                                                                                                                                                                                 derived from intergenic regions. The chimeric promoters are
                                                                                                                                                                                                                                                                                                                                                               Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                        sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-MAR-2000;
             138
                                          63
                                                                    78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                    present sequence represents a chimeric promoter of the invention specification describes chimeric expression promoters. These
                                                                                                                                                                                    producing
GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG 197
                                        eric expression promoter for transgenic plant production, comprisence from promoter comprising vascular expression region replaced sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-647238/62
                                                                                                153;
                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                             Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gruber V,
                                                                                                  Conservative
                                                                                                                                                          BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                               82;
                                                                                                                                                                                    transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99FR-0003925
                                                                                                                                                        98 A; 54 C;
                                                                                                                                                                                                                                                                                                                                                               91pp; English.
                                                                                                              23.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Theisen
                                                                                                                                                                                     plants.
                                                                                                  0;
                                                                                                              Score 140.8;
Pred. No. 3.6
                                                                                                                                                          74 G;
                                                                                                Mismatches
                                                                                                                                                          75 T; 0 other;
                                                                                                                .6e-33;
                                                                                                                            DB 21;
                                                                                                  Indels
                                                                                                                             Length
                                                                                                                             301;
                                                                                                1;
                                                                                                                                                                                                    useful
                                                                                                                                                                                                                                                                                                                                                                                                                                     comprises
                                                                                                Gaps
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ARESULT 13
AAV14019
ID AAV14
AC AAV14
AC AAV14
AC AAV14
XX 18-JU
DT 18-JU
DE CSVMV
XX CASSA
KW L15SU
OS CASSA
XX W0974
PN W0974
PD 24-DE
XX 20-JU
PR 20-JU
PR 20-JU
PA (SCRI
                                                                                                                                                                                                                                                                                                                                                                        밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            망
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                   Cassava vein
                                                                                                                                                                           Cassava vein mosaic virus; CsVMV; promoter; cultivated crop; tissue-specific expression control; transgenic plant; ss.
                                         20-JUN-1996;
                                                                 20-JUN-1997;
                                                                                            24 - DEC - 1997
                                                                                                                                                                                                                                               18-JUN-1998
                                                                                                                                                                                                                                                                                                  AAV14019 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rance I,
              (SCRI ) SCRIPPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric promoter of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                     279 GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                      138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ived from intergenic regions. The chimeric promoters are useful producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                    GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC
                                                                                                                                                                                                                                                                                                                                                                                                                           GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                            GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
                                                                                                                                                                                                                    promoter CVP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>ن</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP;
                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                  mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                82; 91pp;
                                         96US-0020129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99FR-0003925
                                                                 97WO-US10376
              RES INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        128
                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 80 C; 93 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Theisen M;
                                                                                                                                                                                                                                                                                                    392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 140.8;
Pred. No. 4e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                No. 4e-33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ۲,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                    197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               219
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RESULT 14
AAV14021
ID AAV14
XX AAV14
XX AAV14
XX IB-JU
DT 18-JU
XX Cassa
KW Cassa
KW Ctssu
XX Cassa
PT Cassa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents a cassava vein mosaic virus promoter, and is a nucleic acid molecule of the invention. The promoter is capable of initiating transcription of an operably linked heterologous nucleic acid sequence in a plant cell. The CsVMV promoters are active in both monocot and dicot plant species, and therefore can be readily applied to a variety of cultivated crops. Although generally constitutive, the derivative promoters include promoters that can regulate expression in a tissue-specific manner, and therefore are useful for controlling expression of heterologous genes in a tissue-specific manner. The promoters can be used for producing transgenic plants with an altered
            Cassava vein mosaic virus promoter - used to express heterologous DNA sequences for producing transgenic plants having altered
                                                                                                                                                                                                                                                                    Cassava vein mosaic virus; CsVMV; promoter; cultivated tissue-specific expression control; transgenic plant; s
                                                                                                                                                                                                                                           Cassava vein
                                                                                                                                                                                                                                                                                                            CSVMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cassava vein
                                                    WPI; 1998-063157/06
                                                                                                                                  20-JUN-1996;
                                                                                                                                                           20-JUN-1997;
                                                                                                                                                                                       24 - DEC - 1997
                                                                                                                                                                                                                W09748819-A1
                                                                                                                                                                                                                                                                                                                                                                 AAV14021;
                                                                                                                                                                                                                                                                                                                                                                                            AAV14021 standard; DNA; 411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 392 BP; 154 A; 64 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Beachy RN,
                                                                                                       (SCRI ) SCRIPPS RES INST
                                                                                                                                                                                                                                                                                                                                       18-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    143
                                                                                                                                                                                                                                                                                                                                                                                                                                                            263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATCACAAAGGAATCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATGCATTTGTAATTT
                                                                                                                                                                                                                                                                                                            promoter pB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Page 74; 115pp;
                                                                               RN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                               De
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         De Kochko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                           mosaic virus.
                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for producing
                                                                             Kochko A,
                                                                                                                                 96US-0020129
                                                                                                                                                             97WO-US10376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      227
                                                                                                                                                                                                                                                                                                                                                                                                                                                            277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
                                                                               Fauquet C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     promoter -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fauquet C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 120.6; Db ...
                                                                                                                                                                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       romoter - used to express transgenic plants having
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           83 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91 T; 0 other;
                                                                               Verdaguer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Verdaguer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       heterologous
altered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        212
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RESULT 15
AAA96836
ID AAA96
XX AAA96
XX Promo
XX Promo
KW Promo
KW Plant
KW Chime
KW plant
XX Cassa
XX WO200
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Best Local Similarity
Matches 126; Conserv
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                                                                           Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                        Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 411
                                                                                                                                                   WPI; 2000-647238/62.
                                                                                                                                                                                                                                                                                                                                                           WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                           Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                            transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Promoter from intergenic region of Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA96836 standard;
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                                                                                                                                                                                                                                                                                        29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                       (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                         29-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93 GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAAGATAAGGTCGGTGAT 152
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                                                                                                                                                                                   Gruber V,
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Pred. No. 6.3e-27;
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Search completed: April 14, Job time: 196.814 secs

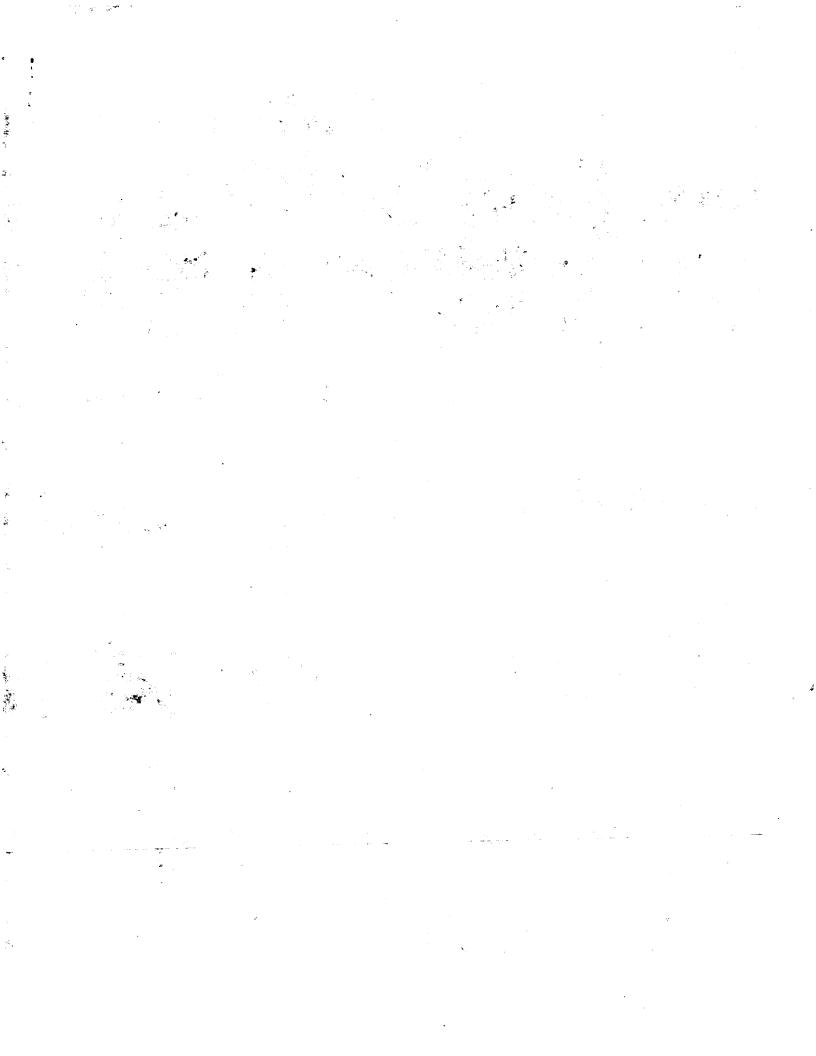
2003, 08:19:41

Claim 4; Page 80; 91pp; English

present sequence represents a promoter fragment from the intergenic

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                             region of Cassava vein mosaic virus. The promoter is used to construct chimmeric expression promoters. These chimmeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                  Sequence 515 BP; 198 A; 79 C; 109 G; 129 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                             The chimeric promoters are useful for producing transgenic plants
336
                                           213
                                                                                                                                                                         216 GAATCTTGAAGACGTAAGCACTGACGACAATGAAAAGAAGAAGATAAGGTCGGTGAT 275
                                                                                                                                                                                                93 GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT 152
                                                                                                           TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGGAAAGTAACCTT 212
ATCACAAAGGAATCT 350
                                           ATGCATTTGTAATTT 227
                                                                                      TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCCGGAAAGTAACCTT 335
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93.3%;
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Pred. No. 6.9e-27;
0; Mismatches 9
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Run on:

Sequence 3 Patent No. Sequence

30, Appl 30, Appl 5254799

Sequence 46, Appl Sequence 35, Appli Sequence 8, Appli Sequence 9, Appli Sequence 46, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 5, Appli

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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600
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
               GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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US-08-247-809A-3
US-08-711-728-3
US-08-764-100-23
US-09-042-426-1
US-09-320-760-1
US-09-320-737-1
US-09-320-737-1
US-09-320-737-1
US-09-320-737-1
US-09-320-737-5
US-09-320-737-7
US-09-320-737-7
US-09-320-737-7
US-09-320-737-7
US-09-320-737-7
US-09-320-736-7
US-09-320-736-7
US-08-450-834-5
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                      Sequence 14, Appl
Sequence 3, Appli
Sequence 3, Appli
Sequence 1, Appli
Sequence 5, Appli
Sequence 1, Appli
                        Sequence
Patent No.
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 Sequence
               Sequence
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US-08-232-463-14
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; Patent No. 5670367
; GENERAL INFORMATION:
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TELEX: 899149

INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
FRIGHT: 7218 base pairs
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Query Match
                                                                                                                                                                                                            FILING DATE:
APPLICATION NUMBER: EP 91
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,76
                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 30
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: DORNER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
ZIP: 223
                                                                 TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
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CITY: Alexandria
STATE: VA
                                                                                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION: 435
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22313-0299
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1800 Diagonal Road,
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US-09-185-244-2
US-09-471-913-6
US-08-894-440-2
US-09-0458-093-2
US-09-065-999-6
US-09-065-999-6
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US-08-729-601A-43
US-09-363-970-35
US-09-011-151-8
US-09-011-151-9
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US-08-463-308-30
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US-08-711-728-5
Score 54.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/08247809A Patent No. 5569823 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 6.38 Matches 26; Conservative
                                                                                                                                                                                                                           OPERATING SYSTEM: DOS
SOPTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/2
FILING DATE: May 23, 1994
CLASSIFICATION: 800
                                              INFORMATION FOR SEQ ID NO:
                                                                                       REGISTRATION NUMBER: 33,141
REFERENCE/DOCKET NUMBER: BA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 332-1700
SEQUENCE CHARACTERISTICS:
LENGTH: 439 base pairs
TYPE: nucleic acid
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                                                                                                                                                    FILING DATE: May 28, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Kurt G. Briscoe
                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Peter H. Schreier; Klaus Stenzel; Gunter Adam; APPLICANT: Edgar Maiss
TITLE OF INVENTION: DEOXYRIBONUCLEIC ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1362
                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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                                                                           (914) 332-1844
                                                                                                                                                                                                                                                                                                                         NEC Powermate 1 Plus
                                                                                                                                                                                                 P 43 178 45.6 (Germany)
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16; Mismatches 168;
                                                                                                                       Bayer
                                                                                                                         9049-KGB
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; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-247-809A-3
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                                                           Best
                                                                         Query Match
                                                  Matches
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                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                           FILING DATE: 23-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 4:
FILING DATE: 28-MAY-1993
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 1 FILING DATE: 03-SEPT CLASSIFICATION: 800 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Peter H. Schreier; Klaus Stenze.
APPLICANT: EGGAR Maiss
TITLE OF INVENTION: DEOXYRIBONUCLEIC ACIDS
NUMBER OF SEQUENCES: 18
TYPE: nucleion STRANDEDNESS:
                                                                                                                                                                                            TELEX:
                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
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                                                            Local
                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: WordPerfect 5.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Tarrytown
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                                                                                                                                                         LENGTH:
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5. 5973135
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                                                           Similarity
                                                                                                                                         nucleic acid
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                                                 Conservative
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                                                           6.7%;
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                                                           Score 40.4; DB 2
Pred. No. 0.0028;
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                                               Mismatches
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                                                                      DB 2;
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                                                                      Length 439;
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RESULT

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US-09-042-426-1
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Best Local Similarity
                                                                                                                                                                               Sequence 1, Application US/09042426
Patent No. 6114608
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 17-MAR-1993
APPLICATION NUMBER: GB 9206016.9
FILING DATE: 19-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO. 57737007is, Allen E.
NAME: NO. 57737007is, 34.190
REGISTRATION NUMBER: 34.490
REFERENCE/DOCKET NUMBER: 137-1061
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 446 base pairs
TYPE: nucleic acid
                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6114608artis Corporation
                                                                                                          TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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TITLE OF INVENTION: Compounds
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
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APPLICANT: De Haan, Petrus T
APPLICANT: Gielen L., Johanne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 354-3592
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                                                                                                                                                                                                                                                                                                                      396 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 445
                                                                                                                                                                                                                                                                                                  274 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 323
STATE: N
         CITY: Summit
STATE: New Jersey
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                                                   STREET:
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                                                 564 Morris Avenue
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Goldbach, Rc
                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                              Irvin J.
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                                                                                                                            Mettler, Paul S. Dietrich, Ralph Sinibaldi
DNA Construct Containing Bacillus
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Pred. No. 0.0028;
                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 446;
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Best Local :
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NAME: HOXie, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/POCKET NUMBER: 135/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (919) 541-861 INFORMATION FOR SEQ ID NO:
FILING DATE: MARCH 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541.8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
VOLFORTY
                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
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HYPOTHETICAL:
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                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/0 FILING DATE: March 13, 1998 CLASSIFICATION: 800
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                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                         COUNTRY: UZIP: 07901
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                                                                                                                         APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                      : Summit
E: New Jersey
TRY: USA
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/ENTION: DNA Construct Containing Bacillus
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                    Floppy disk
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                                                                                                          09/042,426
L3, 1998
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                                      135/1
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                                                             US-09-330-760-1
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               Query Match
Best Local Similarity
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   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6229075artis Corporation
                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS LENGTH: 532 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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CLONE: 35S Promoter
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                                                                                         IMMEDIATE SOURCE:
                                                                                                                     MOLECULE TYPE: DI
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 09/042,426
FILING DATE: MARCH 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                          ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: doub
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: New J
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                                                                            CLONE:
                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/330,760 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER:
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STRANDEDNESS: double
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                                                                            35S Promoter
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                                                                                                                                                                                                 532 base pairs
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DNA Construct Containing Bacillus
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               Score 40.4; DB Pred. No. 0.003;
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                            DB 4;
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                            Length 532;
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US-09-328-473-1
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                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                           Best
                                                GENERAL INFORMATION:
APPLICANT: Irvin J. Mettler, Paul
TITLE OF INVENTION: DNA CONSTRUCT
NUMBER OF SEQUENCES: 11
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: 09/042,426
APPLICATION UNMBER: 09/042,426
FILING DATE: March 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION UNMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
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STREET: J.
CITY: Summit
CITY: New Jersey
I'SA
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CLONE: 35S Promoter
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TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6232533artis Corporation
                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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ADDRESSEE:
STREET: 56
                                                                                                                                                                                                                                                                                                       Local
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                                                                                                                                                                                                                                                                                                      Similarity
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E: No. 6232534artis Corporation 564 Morris Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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DNA Construct Containing Bacillus
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                                                                  S. Dietrich, Ra
Containing Baci
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MEDIUM TYPE: Floppy

COUNTRY:

07901

STATE: New Jersey

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:

APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (919)
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SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/042,426
FILING DATE: March 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX16, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
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REFERENCE/DOCKET NUMBER: 13:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             327 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          274 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 323
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/329,169

FILING DATE: 09-Jun-1999

CLASSIFICATION: <Unknown>
APPLICATION NUMBER: 09/042,426 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                         PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
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STATE: New Jersey
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US-09-330-714A-1
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                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 09/042,426
FILING DATE: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
NAME: HOXIe, Thomas
REGISTRATION NUMBER: 32,993
                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/330,714A
FILING DATE: 11-Jun-1999
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 07901
COMPUTER READABLE FORM:
                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
TYPE: nucleic acid
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ANTI-SENSE: NO
IMMEDIATE SOURCE:
                                                    TOPOLOGY: linear MOLECULE TYPE: DNA (9
                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 135/1 TELECOMMUNICATION INFORMATION:
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                                    HYPOTHETICAL:
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TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/09330714A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 532 base pairs TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                         TELEPHONE: (919)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 564
CITY: Summit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (919) 541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 32,993 REFERENCE/DOCKET NUMBER: 13
                                                                                               STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: No. 6342660artis Corporation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            564 Morris Avenue
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88.0%;
                                                                                                                                                                                                              (919) 541-8614
                                                        (genomic)
                                                                                                                                                                                              541-8689
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CLONE: 35S Promoter; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-330-714A-1
                 RESULT 13
US-09-042-426-5
                                                                                                                                                                                                                                            US-09-328-826-1
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US-09-328-826-1
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Patent No. 6399860
Patent INFORMATION:
GENERAL INFORMATION:
APPLICANT: ITVIN J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 6.7%;
Best Local Similarity 88.0%;
Matches 44; Conservative
Sequence
                                                                                                                                                                    Matches
                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPHONE: (919) 541-8689
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                              327 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/042,426
FILING DATE: MATCH 13, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                          CLONE: 35S Promoter SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                         GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 323
5, Application US/09042426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/328,826 FILING DATE: 09-Jun-1999 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 532 base pairs
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                                                                                                                                                                  Conservative
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                                                                                                                                                                                   6.7%;
                                                                                                                                                                                 Score 40.4;
Pred. No. 0.
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                                                                                                                                                                                   DB 4; Length 532;
.003;
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US-09-291-238-5
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ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPAX: (919) 541-8669
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                       Sequence 5, Application US/09291238
Patent No. 6222104
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 6114608
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HYPOTHETICAL: N
ANTI-SENSE: NO
SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/291,238
                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6222104art1s Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6114608artis Corporation
                                                                                                                                                                                                                                                                                     APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 322 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   327 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 376
                                                                                                                                             CITY: Summit
STATE: New Jersey
COUNTRY: USA
                                                                                                                                   COUNTRY: UZIP: 07901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/0 FILING DATE: March 13, 1998 CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                             STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
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New Jersey
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                                                                                                                                                                                                             564 Morris Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6.7%;
88.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US/09/042,426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 40.4; DB 3; Pred. No. 0.0031;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 560;
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PRIOR APPLICATION DATA:

09/042,426

1998

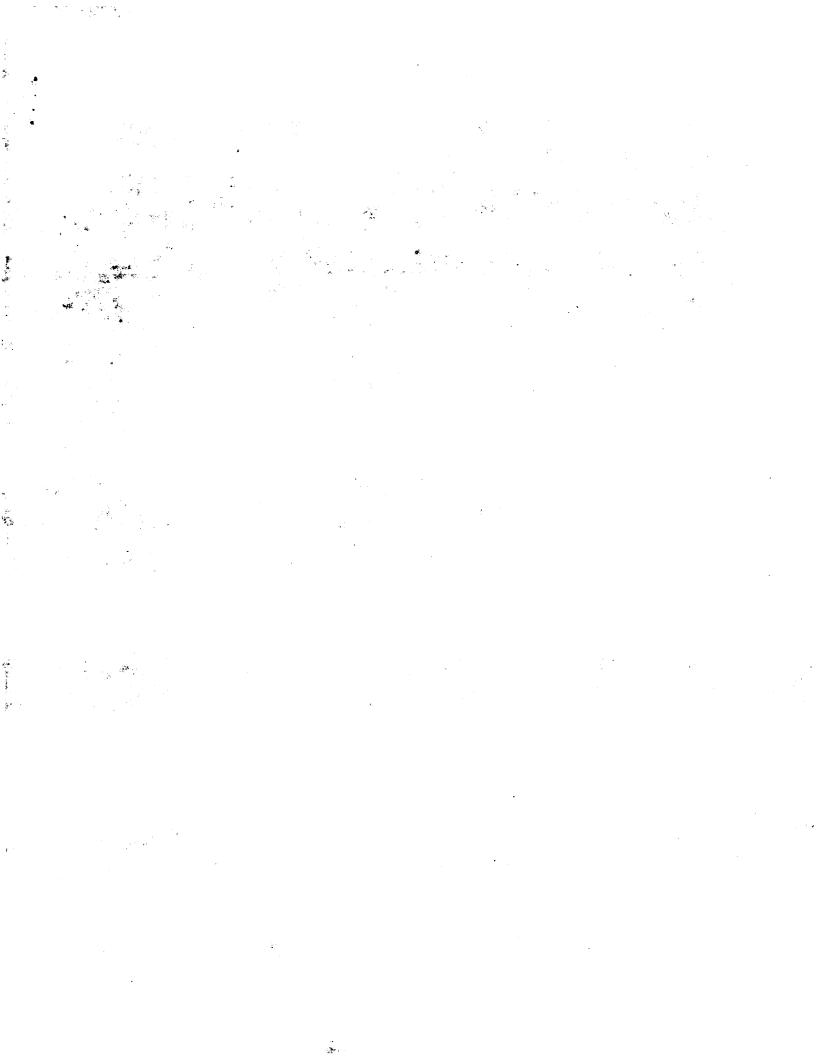
CLASSIFICATION:

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RESULT 15
US-09-330-760-5
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                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/042,420
APPLICATION NUMBER: 13,198
FILING DATE: MATCH 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOXIE, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPAX: (919) 541-8614
TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 560 been saits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 6.7%;
Best Local Similarity 88.0%;
Matches 44; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (919) 541-8689 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS: ADDRESSE: No. 6229075artis Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Irvin J. TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IMMEDIATE SOURCE:
CLONE: 35S Promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
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REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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FILING DATE: March 13, 19
ATTORNEY/AGENT INFORMATION:
NAME: Hoxle, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Summit
STATE: New Jersey
COUNTRY: USA
ZIP: 07901
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    322 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: doi TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                    ENGTH:
                            H: 560 base pairs nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           564 Morris Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
VENTION: DNA Construct Containing Bacillus
EQUENCES: 11
linear
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                double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6;
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                                                                                           ; CLONE: 35S Promoter US-09-330-760-5
                                            Query Match 6.7%;
Best Local Similarity 88.0%;
Matches 44; Conservative
                                                                                                                               MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                 IMMEDIATE SOURCE:
DNA (genomic)
                                            Score 40.4; DB 4;
Pred. No. 0.0031;
0; Mismatches 6;
                                                                  Length 560;
                                              Indels
                                              0;
                                             Gaps
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Search completed: April 14, 2003, 10:28:42 Job time: 67.8557 secs



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Total number of hits satisfying chosen parameters:
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Listing first 45 summaries
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: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
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: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
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US-09-963-803-25

US-09-963-803-2

US-09-963-803-2

US-09-963-803-2

US-09-963-803-2

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7293.024 Million cell updates/sec
Sequence 22, Appl
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Sequence 24, Appl
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Sequence 3, Appl
Sequence 3, Appl
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Sequence 5, Appl
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10, 10, vativa TGCAGG	1 Sequ	NUMBER: US/0 1: 2001-09-26 UMMBER: FR 99/ 1999-03-29 UMMBER: FCT IB 2000-10-05 SE: 39 Version 3.1	ion US/09963 30028922A1 Therapuetics Chimeric exp virus and c		219	7129	700 763 3983 4973	65 65 65 65 65 65 65 65 65 65 65 65 65 6	79 60 63 236 236 332
EE: EY: promoter ON: (1)(600) INFORMATION: 803-22 100.0%; Score 600; DB 9; Length 600; al Similarity 100.0%; Pred. No. 2.5e-170; 600; Conservative 0; Mismatches 0; Indels 0; Gaps 0; AAGCTTGCATGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA 60	Sequence omoter M		9/099 9/22A 9/22A Puet Puet Fic e	ALIGNMENTS	10 10	9 9 9 9	9999	בסבספבי	. פפפפתקת
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			originating from commelina yell:		Sequence 4, Appli Sequence 33, Appli Sequence 8, Appli	Sequence 3, Appli Sequence 101, Appl Sequence 10, Appl	Sequence 9, Appli Sequence 9, Appli Sequence 1, Appli Sequence 15, Appli Sequence 25, Appl	Sequence 5, Appli Sequence 8, Appli Sequence 28, Appl Sequence 346, App Sequence 13, Appl Sequence 22, Appl Sequence 23, Appli Sequence 3, Appli	Sequence 14, Appl Sequence 10, Appli Sequence 9, Appli Sequence 12, Appl Sequence 34, Appl Sequence 35, Appl Sequence 36, Appli Sequence 7, Appli

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Sequence 20, Application US/09963803

Publication No. US20030028922A1

GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow material of inventions in the same cassava vein mosaic virus

FILE OF INVENTION: Virus and cassava vein mosaic virus

FILE REFERENCE: 184332042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT APPLICATION NUMBER: FR 99/03925

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR FILING DATE: 1999-03-29

PRIOR APPLICATION UNUMBER: PCT IB00/00370

PRIOR FILING DATE: 2000-10-05

NUMBER OF SEQ ID NOS: 39

SOFTWARE: Patentin version 3.1

SEQ ID NO 20

LENGTH: 462
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                                                                                                                                                                                                                               NAME/KEY: promoter LOCATION: (1)..(462) OTHER INFORMATION:
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OTHER INFORMATION: promoter MPr1163
FEATURE:
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                                                                                                   AAGCTTGCATGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA 60
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                                   GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC 120
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Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 18433042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 23
LENGTH: 604
TYPE: DNA
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Best Local Similarity
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PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
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                                                                                                                                                GATAAGGTCGGTGATTGTGAAAGAGACATAGAGGGACACATGTAAGGTGGAAAATGTAAGG
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                                                                                               GATAAGGTCGGTGATTGTG-AAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGG
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84.5%;
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Pred. No. 6.1e-84;
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CURRENT APPLICATION UNMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR EILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 19
LENGTH: 393
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US-09-963-803-19
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Best Local
                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: promoter LOCATION: (1)..(393) OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
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ACGACTAGTGATTGATGTGATATCAAGATTGATGT
                                                             GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
                                                                                                                      AGGTGGAAAATGTAAGGGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGACTAGT
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                                             GATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
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US-09-963-803-21
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         Sequence 21, Application US/09963803

Publication No. US20030028922A1

GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 18433042

CURRENT APPLICATION NUMBER: US/09/963,803

CURRENT FILING DATE: 2001-09-26

CURRENT FILING DATE: 2001-09-26
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Best Local Similarity
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TITLE OF INVENTION: Chimeric expression promoters originating:
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05 NUMBER OF SEQ ID NOS: 39 SOFTWARE: PatentIn version 3.1
PRIOR APPLICATION NUMBER: FR 99/03925
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TYPE: DNA
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APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from the promoters of the promoters of the promoters of the price partial parts: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PCT IB00/00370
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NAME/KEY: promoter

LOCATION: (1)..(392)

OTHER INFORMATION:
US-09-963-803-21
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                                                                 FEATURE:
NAME/KEY: promoter
LOCATION: (1)..(317)
OTHER INFORMATION:
1S-09-963-803-3
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PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 21
LENGTH: 392
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Best Local Similarity
Matches 239; Conserv
Query Match
Best Local Similarity
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TYPE: DNA
ORGANISM: Artificial Sequence
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                FEATURE: OTHER INFORMATION: Promoter MPr1116
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96.8%;
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Pred. No. 1.3e-60;
""" matches 8;
Score
Pred.
212.4;
No. 4e-
  4e-54;
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US-09-963-803-25

; Sequence 25, Application US/09963803

; Publication No. US20030028922A1
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; LOCATION: (1)..(472)
; OTHER INFORMATION:
US-09-963-803-25
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APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
TITLE OF INVENTION: virus and cassava vein mosaic virus
TILLE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PT 1B00/00370
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
NUMBER OF SEQ ID NOS: 39
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SEQ ID NO 25
LENGTH: 472
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
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399
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            TGATATCAAGATTGATGT 335
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GGTTCCCAAGCTTTATTT
                                                    TAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTAAGGTCGAAAATGTAAGGGC 197
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91.5%;
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Pred. No. 4.9e-54;
0; Mismatches 21
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RESULT 10
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                                                                                                                                                   GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow |

TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow |

TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow |

TITLE OF INVENTION: Chimeric expression mosaic virus

TITLE OF INVENTION: Chimeric expression mosaic virus

TITLE OF INVENTION: 184332042

CURRENT FILING DATE: 2001-09-26

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR APPLICATION NUMBER: FR 99/03925

PRIOR APPLICATION NUMBER: PCT IB00/00370

PRIOR FILING DATE: 2000-10-05

NUMBER OF SEQ ID NOS: 39

SOFTWARE: Patentin version 3.1

SEQ ID NO 5

LENGTH: 371
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SEQ ID NO 4
LENCTH: 348
TYPE: DNA
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Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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                                                                   ORGANISM: Artificial Sequence
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OTHER INFORMATION: promoter MPr1117
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                                                                                                                              TYPE: DNA
OTHER INFORMATION: promoter MPr1146
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Pred. No. 1.7e-50;
0; Mismatches 1;
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; FEATURE:
; NAME/REY: promoter
; LOCATION: (1)..(301)
; OTHER INFORMATION:
US-09-963-803-7
RESULT 12
US-09-963-803-6
; Sequence 6, Application US/09963803
; Publication No. US20030028922A1
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Best Local S
Matches 153
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PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 7, Application US/09963803 Publication No. US20030028922A1
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE: OTHER INFORMATION: promoter MPr1154
                                                                                                          198 GCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC 233
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                                                                                                                                                                                                                                                63 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAA 122
                                                                                                                                                                                                                                                                                                                     Local Similarity
nes 153; Conserv
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98.7%;
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Pred. No. 1.2e-32;
0; Mismatches 2;
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Pred. No. 3.4e-36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/09963803 Publication No. US20030028922A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                     SEQ ID NO 2
LENGTH: 515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 23.5%;
Best Local Similarity 98.1%;
Matches 153; Conservative
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APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-99-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PCT IB00/00370
Query Match
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                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR ETLING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
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NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                        NAME/KEY: promoter LOCATION: (1)..(51: OTHER INFORMATION:
                                                                                                                                      OTHER INFORMATION:
OTHER INFORMATION:
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                      FEATURE
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OTHER INFORMATION: promoter MPr1147
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Pred. No. 1.4e-32;
0; Mismatches 2
  Score 120.6;
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                                                                                                                                                        the intergenic region of Cassava Vein Mosaic virus
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Length 515;
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PRIOR APPLICATION NUMBER: US 60/177,468
PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 75
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 532
TYPE: DNA
PRESENTATION: Artificial Sequence
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US-09-765-555-1
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GENERAL INFORMATION:
APPLICANT: AGRINOMICS, LLC.
APPLICANT: AGRINOMICS, LLC.
TITLE OF INVENTION: IDENTIFICATION AND CHARACTERIZATION OF A PAGODA PHENOTYPE
FILE REFERENCE: PAGODA
CURRENT FILING DATE: 2001-05-01
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Sin
Matches 126;
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Publication No. US20030037355A1
                                                                                                                                                    Sequence 4, Application US/09847057 Patent No. US20020004943A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 09/620,897 PRIOR FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/765,555
CURRENT FILING DATE: 2002-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: The Scripps Research Institute TITLE OF INVENTION: Methods and compositic TITLE OF INVENTION: expression in plants FILE REFERENCE: 27801-20014.40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                           153 TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT 212
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Pred. No. 1.8e-26;
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SEQ ID NO 4 LENGTH: 8340

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Maximum Match 100%
Listing first 45 summaries
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/cgn2_6/ptodata/1/pna/US080_COMB.seq:*
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44. /cgn2_6/ptodata/1/pna/US6002_COMB.seq: *
46. /cgn2_6/ptodata/1/pna/US6002_COMB.seq: *
47. /cgn2_6/ptodata/1/pna/US6003_COMB.seq: *
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49. /cgn2_6/ptodata/1/pna/US6006_COMB.seq: *
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51. /cgn2_6/ptodata/1/pna/US6006_COMB.seq: *
52. /cgn2_6/ptodata/1/pna/US6009_COMB.seq: *
53. /cgn2_6/ptodata/1/pna/US6009_COMB.seq: *
54. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
55. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
56. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
57. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
58. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
60. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
61. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
62. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
63. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
64. /cgn2_6/ptodata/1/pna/US6011_COMB.seq: *
65. /cgn2_6/ptodata/1/pna/US6012_COMB.seq: *
68. /cgn2_6/ptodata/1/pna/US6012_COMB.seq: *
69. /cgn2_6/ptodata/1/pna/US6021_COMB.seq: *
69. /cgn2_6/ptodata/1/pna/US6022_COMB.seq: *
71. /cgn2_6/ptodata/1/pna/US6022_COMB.seq: *
72. /cgn2_6/ptodata/1/pna/US6022_COMB.seq: *
73. /cgn2_6/ptodata/1/pna/US6023_COMB.seq: *
74. /cgn2_6/ptodata/1/pna/US6023_COMB.seq: *
75. /cgn2_6/ptodata/1/pna/US6023_COMB.seq: *
76. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
77. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
78. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
79. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
79. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
79. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
79. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
80. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
81. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
82. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
83. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
84. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
85. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
85. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
85. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
85. /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
86. /cgn2_6/ptodata/1
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

21	20	19	18	17	16	15	14	13	12	11	10	9	6 0	7	6	5	4	ω	2	₩.	No.	Result
120.6	120.6	120.6	120.6	120.6	120.6	120.6	120.6	120.6	140.8	140.8	152.8	200.4		212.4		281.4	304.6	312.2	373.6	600	Score	
20.1	20.1	20.1	20.1	20.1	20.1	20.1	20.1	20.1	23.5	23.5	25.5	33.4	35.4	35.4	39.0	46.9	50.8	52.0	62.3	100.0	Match I	Query
515	411	411	411	411	392	392	392	392	398	301	371	348	472	317	392	541	393	604	462	600	Match Length DB	
25	16	16	_	1	16	16	۲	_	36	36	36	36	36	36	36	36	36	36	36	36	;	
US-09-641-466-1	US-09-202-838A-4	US-09-202-838-4	PCT-US97-10376A-4	PCT-US97-10376-4	US-09-202-838A-1	US-09-202-838-1	PCT-US97-10376A-1	PCT-US97-10376-1	US-09-963-803-6	US-09-963-803-7	US-09-963-803-5	US-09-963-803-4	US-09-963-803-25	US-09-963-803-3	US-09-963-803-21	US-09-963-803-24	US-09-963-803-19	US-09-963-803-23	US-09-963-803-20	US-09-963-803-22	ID	
Sequence 1, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 6, Appli	Sequence 7, Appli	Sequence 5, Appli	Sequence 4, Appli	Sequence 25, Appl	Sequence 3, Appli	Sequence 21, Appl	Sequence 24, Appl	Sequence 19, Appl	Sequence 23, Appl	Sequence 20, Appl	Sequence 22, Appl	Description	

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Sequence 22, Application US/09963803

GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

TITLE OF INVENTION: Chimeric expression promoters originating from the promoters originating from the promoters originating from the promoters of the 
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US-09-963-803-22
; Sequence 22, Ap)
; GENERAL INFORMA
; APPLICANT: Mer
; TITLE OF INVEN
; TITLE REFERENCE
; FILE REFERENCE
; CURRENT APPLIC
; CURRENT FILING
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Best Local Similarity
Matches 600; Conserv
                                                                                                                                                                                                                                                                                        ORGANISM: Artificial (FEATURE:
OTHER INFORMATION: proFEATURE:
                                                                                                                                                                                                                                              NAME/KEY: promoter LOCATION: (1)..(600) OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAGCTTGCATGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA
             AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA 180
                                                                      GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC
AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
                                                       GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC
                                                                                                                                                                        100.0%; llarity 100.0%; Conservative 0.
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PCT-US97-10376-2
ECT-US97-10376-2
ECT-US97-10376-3
ECT-US97-10376-3
PCT-US97-10376-3
PCT-US97-10376-3
PCT-US97-10376-3
US-09-673-274A-19
EUS-09-673-274A-19
EUS-09-673-274A-19
EUS-09-673-274A-20
US-09-673-274A-20
US-09-673-274A-19
EUS-09-673-274A-20
US-09-673-274A-19
EUS-09-673-274A-20
US-09-673-274B-19
EUS-09-74-926-4
US-09-74-926-4
US-09-74-9388-5
EUS-09-202-838A-5
PCT-US97-10376A-9
                                                                                                                                                                        0;
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Pred. No. 8.
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                                                                                                                                                                         Mismatches
                                                                                                                                                                                    DB 36;
}.9e-163;
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NAME/KEY: promoter

LOCATION: (1)..(462)

OTHER INFORMATION:
US-09-963-803-20
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US-09-963-803-20
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                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 20
LENGTH: 462
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20, Application US/09963803
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                       Query Match
Best Local Similarity
Matches 385; Conserv
                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: promoter MPr1163
                                                                                                                                                                                                                                                                                                                   FEATURE:
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121
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                                                                                                            AAGCTTGCATGCCTGCAGCACTAGTATCCGCCGTCATCAATGACATCATCACAGTACTGA
           AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGAGACATAGAGGACACATGTA
                                                                   GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC
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AACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACATGTA
                                                       GGAGATGAATAGCTAGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGAC
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                                                                                                                                                                       Conservative
                                                                                                                                                                                    62.3%;
95.3%;
                                                                                                                                                                     Score 373.6;
Pred. No. 2.9e
0; Mismatches
                                                                                                                                                                     0;
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2.9e-97;
hes 19;
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                                                                                                                                                                                                 Length 462;
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FEATURE:

NAME/KEY: promoter

LOCATION: (1)..(604)

OTHER INFORMATION:
US-09-963-803-23
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: F99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 23
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity Matches 442; Conserv
                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -09-963-803-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE: OTHER INFORMATION: promoter MPr1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 604
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                                                                                                                                                                                                                                                                                                                                78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                          TGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTAG
                                                                                                                                                                                                     ACGCATGCCACGACTAGTGATTGATGTGATATCAAGATTGATGT 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACGACTAGTGATTGATGTGATATCAAGATTGATGTGATATCTCCACTGACGTAAGGGATG
                                                                                                                                                                                       GCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC------
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                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                         52.0%;
                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                         Score 312.2;
Pred. No. 1.9
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                                                                                                                                  -----CTAGTGATTGATG
                                                                                                                                                                                                                                                                                                                                                                         ; DB 36; Length .9e-79;
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RESULT 5
US-09-963-803-24
Ceruence 24, Ap
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LENGTH: 393
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: VITUS and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
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     Application US/09963803
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Pred. No. 2.6e-77;
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from
ITILE OF INVENTION: Virus and cassava vein mosaic virus
ITILE OF INVENTION: Virus and cassava vein mosaic virus
ITILE OF INVENTION: Virus and cassava vein mosaic virus
ITILE OF INVENTION: Virus and cassava vein mosaic virus
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; NUMBER OF SEQ ID NOS: 3
SOFTWARE: PATENTIN VETS
: SEQ ID NO 21
: LENGTH: 392
: TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                              Sequence 21, Application US/09963803
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
FILE REFERENCE: 184332042
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Best Local Similarity
                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                          PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05
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ORGANISM: Artificial Sequence
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; LOCATION: (1)..(317)
; OTHER INFORMATION:
US-09-963-803-3
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 3
LENGTH: 317
                                                                                                                   Query Match
Best Local
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                                                                                                                  Local Similarity
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                                                   GATTGAT 247
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99.58;
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Pred. No. 1.2e-50;
0; Mismatches 1;
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; LOCATION: (1)..(472)
; OTHER INFORMATION:
US-09-963-803-25
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                                                                                      Sequence 4, Application US/09963803
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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LENGTH: 472
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity
Matches 236; Conserv
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTMARE: Patentin version 3.1
TITLE OF INVENTION: Chimeric expression promoters originating TITLE OF INVENTION: virus and cassava vein mosaic virus FILE REFERENCE: 184332042 CURRENT APPLICATION NUMBER: US/09/963,803 CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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91.5%;
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US-09-963-803-5
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chineric expression promoters originating from commelina yell
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
CURRENT FILING DATE: 2001-09-26
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Best Local Similarity
Matches 212; Conserv
                      Query Match
Best Local Similarity
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       Matches
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NAME/KEY: promoter
LOCATION: (1)..(348)
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                          NAME/KEY: promoter LOCATION: (1)..(371) OTHER INFORMATION:
                                                                                                                                                                 OTHER INFORMATION:
                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                FEATURE:
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       154;
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       Conservative
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                      25.5%;
98.7%;
     Score 152.8;
Pred. No. 2.3e
0; Mismatches
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Pred. No. 3.7e-47;
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2.3e-33;
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; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(301)
; OTHER INFORMATION:
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                                                                                                                                                     Sequence 6, Application US/09963803
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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Best Local Similarity
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
SOFTWARE:
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                                                                              CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
                                              PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05
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                                NUMBER OF SEQ ID NOS:
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              PatentIn version 3.1
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Pred. No. 6.4e-30;
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Best Local :
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APPLICATION NUMBER: US 6
FILING DATE: 20-JUN-1996
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Relaction CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                         ANTI-SENSE:
                                                                                                                                                                                                                                                                                    HYPOTHETICAL:
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                213
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                                                                                                                                93 GAATATTGAAGACGTAAGCACTGACCAACAATGAAAAGAAGAAGAAGATAAGGTCGGTGAT 152
                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAA 137
                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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                                                                TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGGCGGAAAGTAACCTT 212
                                                                                                                  GAATCTTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGAAGATAAGGTCGGTGAT
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              ATGCATTTGTAATTT
                                                TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT
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98.1%;
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              227
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Pred. No. 4.9e-24;
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263 ATCACAAAGGAATCT 277

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RESULT 15
US-09-207-838-1
; Sequence 1, Application US/09202838
; GENERAL INFORMATION:
; APPLICANT: Verdaguer, Bertrand
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Best Local
                                                                                                                                                                                                                                                                                                     Matches 126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, \
CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US97/10376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (619) 784-9399 INFORMATION FOR SEQ ID NO: :
                                                                                                                                                                                                                                                                                                                                                                                             TOPULOUS TYPE: DN
                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: I
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TITLE OF INVENTION: CASSAVA VEIN MOSAIC VIRUS PROMOTERS
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: 50
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (619) 784-2937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 20-JUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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ZIP: 920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 10550 North CITY: La Jolla STATE: California
                                                                                                                                                                                                                                                      93 GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAAGAAGATAAGGTCGGTGAT 152
                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
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de Kochko, Alexandre
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Search completed: April 14, 2003, 12:06:34 Job time: 2423.62 secs
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LENGTH: 392
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Best Local Similarity 9...
126; Conservative
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NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/0 PRIOR FILING DATE: 1996-06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT, PRIOR FILING DATE: 1997-06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/202,838
CURRENT FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Fauquet, Claude
TITLE OF INVENTION: CASSAVA VEIN MOSAIC VIRUS PROMOTERS AND USES THEREOF
FILE REFERENCE: MYCOO11S
                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Cassava vein mosaic virus
                                                                                                                                                                               263
                                                                           ATGCATTTGTAATTT 227
                                                                                                                                    TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT 212
                                                          ATCACAAAGGAATCT 277
                                                                                                                      TGTGAAAGAGACATAGAGGACACATGTAAGGTGGAAAATGTAAGGGCGGAAAGTAACCTT
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Beachy, Roger N.
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Pred. No. 4.9e-24;
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Maximum DB seq length: 2000000000
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Perfect score:
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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      Query
Match
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Gapop 10.0 , Gapext 1.0
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600
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    /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
    /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
    /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
    /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
    /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
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    US-09-765-55A-1
US-09-673-274B-29
US-09-673-274B-29
US-10-380-935-30
US-10-380-935-37
US-10-391-414-106-147
US-10-391-416-147
US-10-391-416-147
US-10-130-150-18
US-10-130-150-18
US-10-130-764-4
US-10-160-764-4
US-10-160-764-48
US-10-160-764-49
US-10-160-764-49
US-10-160-764-49
US-10-160-764-49
US-10-160-764-52
US-10-160-764-52
US-10-160-764-90
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US-10-160-764-91
US-10-160-764-91
US-10-160-764-91
US-10-160-764-91
US-10-198-478-16
US-10-198-478-16
US-10-198-478-13
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Sequence 1, Appli
Sequence 20, Appl
Sequence 30, Appl
Sequence 37, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 40, Appl
Sequence 50, Appl
Sequence 40, Appl
Sequence 52, Appl
Sequence 52, Appl
Sequence 52, Appl
Sequence 52, Appl
Sequence 50, Appl
Sequence 14, Appl
Sequence 14, Appl
Sequence 15, Appl
Sequence 16, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 13, Appl
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ch 20.1% 1 Similarity 93.3% 1 Similarity 93.3% 1 Similarity 93.3% 126; Conservative AATATTGAAGACGTAAGCAC	cation US/097 ON: ON: Cripps Resear as, Carlos e, Justin ita, Bipin ON: Methods a ON: expressi 27801-2014.2 ION NUMBER: U ATE: 2001-01 N NUMBER: 09/ E: 2000-07-21 N NUMBER: 09/ E: 2000-01-21 CON: 75 Q for Windows ON: Promoter ON: Promoter	3034 4182 5449 5449 6975 9555 13199 17476 6944	13737 250715 206 382 842 2267 2267 2873 2873 2873 2873 2873 3002 3034
18; 38; H-I-I ACT ACA ACA	US/0976 Researc los in in plots in plot		91999
20.1%; Score 120.6; DB 6; Length 532; y 93.3%; Pred. No. 1.5e-25; y 93.3%; Pred. No. 1.5e-25; Indels 0; G CGTAAGCACTGACGACAACAATGAAAGAAGAAGATAAGGTCGGTGAT			US-09-538-414-10 US-09-949-016-13294 US-10-204-889-9 PCT-US02-17853-22 US-10-321-434-6 PCT-US02-17853-11 PCT-US03-034-35-55 US-10-353-454-38 US-10-353-454-57 PCT-US03-03435-48 US-10-353-03435-48
Length 532; Indels 0; Gaps AAGATAAGGTCGGTGAT 152	,	Sequence 48, Appli Sequence 7, Appli Sequence 57, Appl Sequence 40, Appl Sequence 57, Appl Sequence 6, Appli Sequence 2, Appli Sequence 7, Appli Sequence 7, Appli Sequence 893, App Sequence 113, App	Sequence 10, Appl Sequence 13294, A Sequence 22, Appli Sequence 6, Appli Sequence 6, Appli Sequence 11, Appl Sequence 38, Appl Sequence 55, Appl Sequence 57, Appl Sequence 57, Appl Sequence 57, Appl Sequence 57, Appl
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351 ATCACAAAGGAATCT 365

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APPLICANT: Brookhart, Gary
APPLICANT: Hoffman, Jules
FILE OF INVENTION: GENE CODING FOR HELIOMICINE, AND USE THEREOF
FILE REFERENCE: A33595-PCT-USA (0726667,0166)
CURRENT APPLICATION NUMBER: US/09/673,274B
CURRENT FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: PCT/FF99/00843
PRIOR FILING DATE: 1999-04-12
PRIOR FILING DATE: 1999-04-12
PRIOR FILING DATE: 1999-04-15
PRIOR FILING DATE: 1998-04-15
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 48
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                US-09-673-274B-20 : Sequence 20, Application US/09673274B ; GENERAL INFORMATION:
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PRIOR FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: FR 98 04933
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 48
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19, Application US/09673274B GENERAL INFORMATION:
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TITLE OF INVENTION: GENE CODING FOR HELIOMICINE, AND USE THEREOF
FILE REFERENCE: A33595-PCT-USA (0726667.0166)
CURRENT APPLICATION NUMBER: US/09/673.274B
CURRENT FILING DATE: 1999-04-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Lamberty, Mireille
APPLICANT: Bulet, Phillipe
APPLICANT: Brookhart, Gary
                                                                                                                                                                                                                                                                                                                       APPLICANT: Lamberty, Mireille APPLICANT: Bulet, Phillipe
TYPE: DI
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ORGANISM: Artificial Sequence
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LOCATION: (569)...(832)
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hes 126;
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DNA
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Brookhart, Gary
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Pred. No. 1.8e-25;
Pred. No. 1.8e-25;
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US-10-380-935-37
, Sequence 37, Application US/10380935
; GENERAL INFORMATION:
; APPLICANT: WARNER, Simon, Anthony, James
; APPLICANT: HAWKES, Timothy, Robert
                                                                                                                                                                                                                                                                                        ; FEATURE:
; OTHER INFORMATION: Enhancer
US-10-380-935-30
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; LOCATION: (767)...(1030)
US-09-673-274B-20
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APPLICANT: WARNER, Simon, Anthony, James
APPLICANT: HAWKES, Timothy, Robert
APPLICANT: ANDREWS, Christopher, John
TITLE OF INVENTION: HERBICIDE RESISTANT PLANTS
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Best Local S
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LENGTH: 309
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: GB0027693.1 PRIOR FILING DATE: 2000-11-13 PRIOR APPLICATION NUMBER: GB0023910.3 PRIOR ETLING DATE: 2000-09-29 NUMBER OF SEQ ID NOS: 70 SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/380,935
CURRENT FILING DATE: 2003-03-19
PRIOR APPLICATION NUMBER: GB0023911.1
PRIOR FILING DATE: 2000-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PPD50594/WO
                                                                                                                                                                                                                                                                                                                                       ORGANISM: Cauliflower Mosaic Virus
                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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NAME/KEY: CDS
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                                                                                                                                                242 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 291
                                                                                                                                                                 396 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 445
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Similarity 88.0%;
44; Conservative
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93.3%;
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Pred. No. 0.073;
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PRIOR APPLICATION NUMBER: GB0023911.1
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: GB0027693.1
PRIOR FILING DATE: 2000-11-13
PRIOR APPLICATION NUMBER: GB0023910.3
PRIOR FILING DATE: 2000-09-29
NUMBER OF SED ID NOS: 70
SOFTWARE: PatentIn version 3.1
SEQ ID NO 37
LENGTH: 522
TYPE: DNA
PRIOR FILING: Figwort Mosaic Virus
FERMIDED:
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; Sequence 147, Application US/60434166
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SEQ ID NO 10
LENGTH: 835
TYPE: DNA
ORGANISM: Cauliflower mosaic virus
                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Mendel B
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Best Local
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Stress Induced Promoter Derived From Rice FILE REFERENCE: 382.1041
CURRENT APPLICATION NUMBER: US/10/391,414
CURRENT FILING DATE: 2003-03-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SHINOZAKI, Kazuko
APPLICANT: KATSURA, Koji
APPLICANT: ITO, Yusuke
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TITLE OF INVENTION: HERBICIDE RESISTANT PLANTS
FILE REFERENCE: PPD50594/WO
CURRENT APPLICATION NUMBER: US/10/380,935
CURRENT FILING DATE: 2003-03-19
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                                       APPLICANT
     PPLICANT
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                                                                                                                             Mendel Biotechnology
Creelman, Robert A.
Jiang, Cai-Zhong
Reuber, T. Lynne
Riechmann, Jose Luis
                                                                                                              Creelman, Robe
Haake, Volker
                                                       Heard, Jacqueline E. Adam, Luc J.
                                                                                           Ratcliffe, Oliver
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Pred. No. 0.098;
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; OTHER INFORMATION: G2227 US-60-434-166-147
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CURRENT FILING DATE: 2002-11-22
PRIOR APPLICATION NUMBER: US/09/506,720
PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/120,880
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: 60/121,037
PRIOR FILING DATE: 1999-02-22
PRIOR FILING DATE: 1999-02-22
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SEQ ID NO 147
LENGTH: 1683
                                                                          SOFTWARE: PatentIn
SEQ ID NO 1
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Best Local Similarity
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TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES IN PLANTS
FILE REFERENCE: MBI-0051 P
CURRENT APPLICATION NUMBER: US/60/434,166
CURRENT FILING DATE: 2002-12-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/135,134
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 60/144,153
PRIOR FILING DATE: 1999-07-15
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                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                        PRIOR FILING DATE:
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                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/162,656
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APPLICANT: Creelman, Robert
NITLE OF INVENTION: PLANT GENE SEQUENCES II
FILE REFERENCE: MBI-0007
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            TYPE: DNA
ORGANISM: Arabidopsis thaliana
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FEATURE:
                                                          LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 58
                                                                                                                                                                                                                                                                                                FILING DATE: 1999-03-11
APPLICATION NUMBER: 60/
FILING DATE: 1999-04-15
                                                                                                                                                                              APPLICATION NUMBER: 60/
FILING DATE: 1999-10-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 44; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/10302267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jiang, Cai-Zhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Adam, Luc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Samaha, Raymond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yu, Guo-Liang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reuber, Lynne
Zhang, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pineda, Omaira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fromm, Michael
Heard, Jacqueline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Riechmann, Jose Luis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6.7%;
                                                                                                                                                                                                  60/161,143
                                                                                                                                                                                                                                                                                                                    60/129,450
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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(144)..(2024)

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RESULT 11
US-10-130-150-15
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OTHER INFORMATION: and ER retaining signal peptides
US-10-130-150-18
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Sequence 15, Application GENERAL INFORMATION:
                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 18
LENGTH: 3288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18, Application US/10130150 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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SEQ ID NO 13
LENGTH: 3212
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TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THERBOF
FILE REPERENCE: 02/23831
CURRENT APPLICATION NUMBER: US/10/130,150
CURRENT FILING DATE: 2000-11-15
NUMBER OF SEQ ID NOS: 24
                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Shoseyov, Oded
TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THEREOF
FILE REFERENCE: 02/23831
                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/130,150 CURRENT FILING DATE: 2000-11-15
                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                     ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      197 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                    197 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 246
                                                                                                                                   258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
nes 44; Conserv
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                                                                                                                                                                     Conservative
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88.0%;
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                 US/10130150
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Pred. No. 0.
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                                                                                                                                                                   Mismatches
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                                                                                                GENERAL INFORMATION:
                                                                                                                Sequence 4, Application US/10160764
                                                        APPLICANT: Huang, Yaran APPLICANT: Chalifoux, Maryse
               APPLICANT:
                                                APPLICANT:
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TITLE

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INVENTION:

Angela Patricia Compositions and

and Methods

of

Increasing Stress Tolerance

Wang, Yang Kuzma, Monika Maria Gilley, Angela Patr

3209 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 3258

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; OTHER INFORMATION: Sequence of recombinant construct expressing BGL1 protein fuse; OTHER INFORMATION: signal peptide US-10-130-150-15
                                                                                                                 US-10-160-764-40
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US-10-160-764-40
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Matches
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                                              Matches
                                                             Query Match
Best Local
                                                                                                                                                                                                                               SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                            APPLICANT: Kuzmā, Monika Maria
APPLICANT: Gilley, Angela Patriola
TITLE OF INVENTION: Compositions and Metho
TITLE OF INVENTION: 1n Plants
FILE REFERENCE: 22542-008
CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Shoseyov, Oded
TITLE OF INVENTION: ASPERGILLUS NIGER BETA-GLUCOSIDASE GENE, PROTEIN AND USES THER
FILE REFERENCE: 02/23831
                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/348,909 PRIOR FILING DATE: 2001-10-22
                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 2002-05-31 PRIOR APPLICATION NUMBER: 60/294,766 PRIOR FILING DATE: 2001-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Huang, Yafan APPLICANT: Chalifoux, Maryse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 33
                                                                                                                               OTHER INFORMATION:
                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                              LENGTH: 5247
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial FEATURE:
258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
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                                                             Similarity
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Similarity 88.0%;
44; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wang, Yang
                                                Conservative
                                                                                                                             Description of Artificial Sequence: Plasmid pBII21-35S-AtFTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence
                                                             6.7%;
88.0%;
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                                                            Score 40.4;
Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                             and Methods
                                                Mismatches
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                                                                           Length
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                                                Indels
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                                                                               5247;
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                                              Gaps
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                                              0;
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TITLE OF INVENTION: in Plants

FILE REFERENCE: 22542-008

CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILLING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR FILING DATE: 2001-02
NUMBER OF SEQ ID NOS: 90

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RESULT 15
US-10-160-764-52
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                                                                                                                                                                                                                                                                                                                           US-10-160-764-48
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US-10-160-764-48
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SEQ ID NO 48
LENCTH: 5511
TYPE: DNA
ORGANISM: Artificial Sequence
                                                        Sequence 52, Application US/10160764 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 4
LENGTH: 5250
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local 9
                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                    Query Match
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APPLICANT: Huang, Yafan
APPLICANT: Challfoux, Maryse
APPLICANT: Wang, Yang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/348,909
PRIOR FILING DATE: 2001-10-22
NUMBER OF SEQ ID NOS: 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Gilley, Angela Patricia TITLE OF INVENTION: Compositions and Methods TITLE OF INVENTION: in Plants FILE REFERENCE: 22542-008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Huang, Yafan
APPLICANT: Chalifoux, Maryse
APPLICANT: Wang, Yang
                                                                                                                                                                                                                                                                                                                            FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Plasmid OTHER INFORMATION: pBI121-355-Anti-AtFTB
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                                                                                                                                                                                         258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
                                                                                                                                                                                                                                                               Match 6.7%;
Local Similarity 88.0%;
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Local Similarity 88.0%;
les 44; Conservative
                                                                                                                                                                                                                                               44;
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                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 40.4; DB 9;
Pred. No. 0.17;
0; Mismatches 6;
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                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of Increasing Stress Tolerance
                                                                                                                                                                                                                                                                                9;
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Search completed: April 14, 2003, 12:27:55 Job time: 490.052 secs
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SEQ ID NO 52

LENGTH: 5715

TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Plasmid
OTHER INFORMATION: pBI121-35S-AtFTB
                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                          Matches 44; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kuzma, Monika Maria
APPLICANT: Gilley, Angela Patricla
TITLE OF INVENTION: Compositions and Methods of Increasing Stress Tolerance
TITLE OF INVENTION: in Plants
FILE REFERENCE: 22542-008
CURRENT APPLICATION NUMBER: US/10/160,764
CURRENT FILING DATE: 2002-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/294,766
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/348,909
PRIOR FILING DATE: 2001-10-22
NUMBER OF SEQ ID NOS: 90
                                                                                      3209 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATCCCACTA 3258
                                                                                                             258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 307
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Perfect score:
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BH746858 SALK_0036
BM162517 EST555040
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ALIGNMENTS

	TITLE JOURNAL COMMENT	ORGANISM ORGANISM REFERENCE AUTHORS	RESULT 1 BH746858 LOCUS DEFINITION ACCESSION VERSION KEYWORDS
The Salk Institute Genomic Analysis Laboratory (SIGNAL) The Salk Institute for Biological Studies 10010 N. Torrey Pines Road, La Jolla, CA 92037, USA Tel: 858 453 4100 x1752 Fax: 858 558 6379 Email: ecker@salk.edu This is single pass sequence recovered from the left border of	A Sequence-Indexed Library of Insertion Mutations in the Arabidopsis Genome Unpublished (2001) Contact: Joseph R. Ecker	thale cress. Arabidopsis thaliana Arabidopsis thaliana Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis. 1 (bases 1 to 224) 1 (bases 1 to 224) Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P.,	BH746858 224 bp DNA linear GSS 27-FEB-2002 SALK_003694.51.40.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_003694.51.40.x, DNA seguence. BH746858 BH746858.1 GI:18959973 GSS.

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The Institute for Genomic Research
9712 Medical Center Drive, Rockvill
Tel: 301-530-9319
Fax: 301-838-0208
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Plasmodium yoelii yoelii
Plasmodium yoelia Apicomplexa; Haemosporida; Plasmodium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST
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                                                                                                                                                                                                                                                                                                                                         FOR clone info, please contact the Malaria Research and Reference Reagent Resource Center, ATCC
                                                                                                                                                                                                                                                                                                                                                                              Email: carlton@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Jane Carlton
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/clone="StAL_003694.51.40.x"
/clone="StAL_003694.51.40.x"
/clone="lb="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 68 c 54 g 50 t
/dev_stage="Asexual blood stages"
/lab_host="E. coli XL-1 Blue"
/note="Vector: pAD-GAL4, At 20-25% parasitemia, blood collected from BALB/cByJ mice infected with Py17XL parasites, and leukocytes removed by passage over microcrystalline cellulose collumns. Total RNA was isolated using the guanidinium isothiocyanate method, mRNA isolated using oligo(dT)-cellulose chromatography
                                                                                                                                                                                 /db_xref="taxon:73239"
/clone="PYCKT64"
                                                                                                                                                                                                                     /organism="Plasmodium yoelii yoelii"
/strain="17XL"
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/strain="Columbia 0"
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/clone_lib="RPCI-98" /note="end : TET3"

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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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BACR08K10 of RPCI-98 library f
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Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           First strand cDNA synthesis was completed using a 50-base primer and reverse transcriptase in the presence of 5-methyl dCTP. After second strand synthesis, uneven termini were treated with Pfu DNA polymerase and EcoRI adaptors ligated to the blunt ends. The sample was cleaved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                isolated."
67 c
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                         /organism="Drosophila
/db_xref="taxon:7227"
                                                                                                                   Location/Qualifiers
/clone="BACR08K10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      52.2%;
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rom Drosophila melanogaster (fruit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GACAACAATGAAAAGAAGAAGATAAGGTCGGTGATTGTGAAAGAGACATAGAGGACACAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RAARADRRWTTKGKTTTATWTTWAARAAWWAWWAWATTTATWTTT 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GCCACGACTAGTGATTGATGTGATATCAAGATTGATGTGATATCT 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTARAAWDWWTWKAWDWAKWDWKTRADRWDRWAADTWTDARKADRDWAKARAWRARRDRA 579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTAAGGTGGAAAATGTAAGGGCCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGACT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parasite Genomics Group
The Institute for Genomic |
9712 Medical Center Drive,
Tel: 301-530-9319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Jane Carlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Plasmodium yoelii EST project at TIGR Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Carlton, J.M., Daly T.M., Long, C.A., Bergman, L.W., Fraser, C.M. and Carucci, D.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Plasmodium
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BM161725
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: carlton@tigr.org
For clone info, please contact the
                                                                                                                                                                                                                                                                                                                                                                                               http://www.malaria.mr4.org/mr4pages/index.html
                                                                                                                                                                                                                                                                                                                                                                                                                   Reagent Resource Center,
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                                                                                                                                                                                                                                                                                                                                                                                 primer:
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              /dev_stage="Asexual blood stages"
/lab_host="E. coli XI-1 Blue"
/lab_host="E. coli XI-1 Blue"
/note="Yector: pAD-GAL4; At 20-25% parasitemia, blood was collected from BALB/CByJ mice infected with Py17XL parasites, and leukocytes removed by passage over microcrystalline cellulose collumns. Total RNA was isolated using the guanidinium isothiocyanate method, and mRNA isolated using oligo(dT)-cellulose chromatography. First strand cDNA synthesis was completed using a 50-base primer and reverse transcriptase in the presence of 5-methyl dCTP. After second strand synthesis, uneven termini were treated with Pfu DNA polymerase and EcoRI adaptors ligated to the blunt ends. The sample was cleaved with XhOI and separated on a Sephacryl 5-500 column.
   Size-fractionated cDNA was precipitated and
                                                                                                                                                                                                                                                                                                               /strain="17XL"
                                                                                                                                                                                                                                                                                                                             /organism="Plasmodium
                                                                                                                                                                                                                                                             /clone_lib="PyBS"
                                                                                                                                                                                                                                                                               /clone="PYCKI31"
                                                                                                                                                                                                                                                                                               /db_xref="taxon:73239"
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Query Match
Best Local Similarity
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130 GGCGCAAGCAATCATATTATATTATGCCTTATTATTTATG 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (***).

Contact: Joseph R. Ecker

Salk Institute Genomic Analysis Laboratory (SIGnAL)

The Salk Institute for Biological Studies

The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10010 N. Torrey Pines Road,
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alonso, J.M., Leissé, T.J., Barajas, P., Chen, H., Cheuk, C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Predni, Zimmerman, J. and Ecker, J.R.
A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis Genome Unpublished (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              This is single pass sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                        TDNA tagged
each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html* a 26 c 25 g 30 t
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                                                                                                                                                                                             /clone="SALK_049464.49.75.x"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
                                                                                                                                                                                                                                                                             /organism="Arabidopsis
/strain="Columbia 0"
                                                                                                                                                                    /note="PCR was performed on Arabidopsis thaliana
                                                                                                                                                                                                                                                      /db_xref="taxon:3702"
                                                                                                                                                                                                                                                                                                                                                            ocation/Qualifiers
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53.4%;
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Pred. No.
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Prednis, L., Shinn,
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6.7%; 88.0%;

Score 40.4; Pred. No. 2.

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Length 112,

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RESULT 7
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                                                                                                                                                                                                                    396 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 445
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BH749349
BH749349.1
                                                                 BH619283 153 bp DNA linear GS SALK_040762 Arabidopsis thaliana TDNA insertion lines thaliana genomic clone SALK_040762, DNA sequence.
BH619283
                                    GSS
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute or Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Arabidopsis
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C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
, Zinmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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                 thale cress
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                              /strain="Columbia 0"
/db_xrefa"taxon:3702"
/db_xrefa"taxon:3702"
/clone="ib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.saik.edu/tdna_protocols.html"
a 36 c 30 g 35 t
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Best Local S
Matches 44
                                                                                                                                                 TITLE
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BH747013
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396 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 445
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Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R.,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                               Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                                                                                     Unpublished (2001)
Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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                          Tel: 858 453 4100 x1752 Fax: 858 558 6379
                                                                                                                                                                                                                       Alonso,J.M., Lowers,M., Kim,C.J., runner.
,C., Jeske,A., Karnes,M., Kim,C.J., runner.
, Zimmerman,J. and Ecker,J.R.
, Zimmerman,J. and Ecker,J.R.
, Dimerman,J. and Ecker,J.R.
                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 153)
Alonso, J.M., Leisse, T.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arabidopsis thaliana
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ecker@salk.edu
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note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 34 c 36 g 47 t
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/db_xref="taxon:3702"
/clone="SALK_040762"
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88.0%;
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0; Mismatches
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Prednis,L., Shinn,
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1 (bases 1 to 158)
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                                                                                                                                                                                                                                                                                                                                                                                                                 10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis Genome 
Unpublished (2001)
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C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Predni
Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in
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    QJ
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 37 c 33 g
                                        /clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used ca
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/clone="SALK_045100.51.10.x"
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/strain="Columbia 0"
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/clone="SALK_008070.43.05.x"
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http://signal.salk.edu/tdna_protocols.html" 37 g 49 t
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BH746616 175 bp DNA linear GSS SALK_045268.54.50.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK_045268.54.50.x,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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A. Sequence Indexed Library of Insertion Mutations in
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/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/clone_lib="Arabidopsis thaliana TDNA insertion lines |
/note="PCK was performed on Arabidopsis thaliana lines |
each of which contains one or more TDNA insertion |
elements. The resultant fragment for each line was |
directly sequenced to determine the genomic sequence at |
the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"
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Stanford University
855 California Ave, 1
Tel: 650 723 2227
Fax: 650 725 8221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                       Unpublished (2001 Contact: Walbot V
                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC clade; Panicoldeae; Andropogoneae; Zea.
                                                                                                                                                                                                                                   Zea mays
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GSS.
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                                                                     Department of Biological Sciences
                                                                                                                                        Walbot, V.
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C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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BH746616.1
                                                                                                                         Maize genomic sequences found using
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/clone="SALK_045268.54.50.x"
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/clone="PCR was performed on Arabidopsis thaliana lines notes of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"
a 43 c 40 g 38 t
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∕strain="Columbia 0"
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327 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 190)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,F., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                      Salk Institute Genomic Analysis Laboratory (SIGnAL) The Salk Institute for Biological Studies 10010 N. Torrey Pines Road, La Jolla, CA 92037, USA Tel: 858 453 4100 x1752 Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                            Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
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                                                                                                               Class: TDNA tagged
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                                                                                                                                                                  This is single pass sequence
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/lab_host="DHIOB"
/lab_host="DHIOB"
/note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site_1: BamHI; Site_2: BgIII; RescueMu is a 4.9 kb, modified malze Mu transposon designed to allow plasmid rescue from total genomic DNA. Mu elements insert preferentially into transcription units. For more information on RescueMu, go to the web site www.zmdb.iastate.edu and follow the links for 'RescueMu.' Grid I was grown at Berkeley in 2001. DNA was extracted from leaf punches, double digested using BamHI and BgIII, and ligated to form circular plasmids. DHIOB cells were transformed and then screened on LB plates with ampicillin."
                                                                                                                                                                                               ecker@salk.edu
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/dev_stage="adult"
/lab hoot=""
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/clone_lib="1008 - Rescheme (7-1)
/organism="Arabidopsis thaliana"
/strain="Columbia 0"
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396 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACGACTA 445
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Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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A Sequence-Indexed Library of Insertion Mutations in
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 52 c 41 g 45 t
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                                           Score 40.4; D
Pred. No. 3.1;
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Search completed: April 14, Job time: 1396.64 secs
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                                                                                                                                                                                                                                                                         BASE COUNT
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                                                                                 Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Go,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Zimmerman,J. and Ecker,J.R.

A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1 (bases 1 to 214)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BH747829 214 bp DNA linear GSS 27-SALK_035354.55.75.x Arabidopsis thaliana TDNA insertion lin Arabidopsis thaliana genomic clone SALK_035354.55.75.x, DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Class: TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis Genome
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                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      is single pass sequence recovered from the
                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                         58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ecker@salk.edu
                                                                                                                                                                                                                                                                /clone_lib="Arabidopsis thallana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thallana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 56 c 52 g 48 t
                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="SALK_035354.55.75.x"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /strain="Columbia 0"
/db_xref="taxon:3702"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Arabidopsis thaliana"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                                                                                     6.7%;
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Pred. No. 3.1;
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Prednis,L., Shinn,P.
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